## SEQUENCE LISTING

```
<110> Walke, D. Wade
     Wang, Xiaoming
     Scoville, John
     Turner, C. Alexander Jr.
<120> Novel Human Semaphorin Homologs and Polynucleotides Encoding the Same
<130> LEX-0177-USA
<150> US 60/205,274
<151> 2000-05-18
<150> US 60/208,893
<151> 2000-06-02
<160> 50
<170> FastSEQ for Windows Version 4.0
<210> 1
<211> 3150
<212> DNA
<213> homo sapiens
<400> 1
                                                                      60
atgacqqtqq ttaatccaca agacctgcag ccgtgggtct ctaacttcac ctaccctgga
gcccgggatt tctcccagct ggctttggac ccctccgggr accagctcat cgtgggagcc
                                                                     120
aggaactacc tcttcagact cagccttgcc aatgtctctc ttcttcaggc cacagagtgg
                                                                     180
240
cagaactacg tgcgagtcct gatcgtcgcc ggccggaagg tgttcatgtg tggaaccaat
                                                                     300
                                                                     360
gccttttccc ccatgtgcac cagcagacag gtgggggaacc tcagccggac tattgagaag
atcaatggtg tggcccgctg cccctatgac ccacgccaca actccacagc tgtcatctcc
                                                                     420
tcccaggggg agctctatgc agccacggtc atcgacttct caggtcggga ccctgccatc
                                                                     480
taccgcagcc tgggcagtgg gccaccgctt cgcactgccc aatataactc caagtggctt
                                                                     540
                                                                     600
aatgagccaa acttcgtggc agcctatgat attgggctgt ttgcatactt cttcctgcgg
gagaacgcag tggagcacga ctgtggacgc accgtgtact ctcgcgtggc ccgcgtgtgc
                                                                     660
aagaatgacg tggggggccg attcctgctg gaggacacat ggaccacatt catgaaggcc
                                                                     720
                                                                     780
cggctcaact gctcccgccc gggcgaggtc cccttctact ataacgagct gcagagtgcc
                                                                     840
ttccacttgc crgagcagga cctcatctat ggagttttca caaccaacgt aaacagcaty
geggettetg etgtetgege etteaacete agtgetatet eccaggettt caatggeeca
                                                                     900
                                                                     960
tttcgctacc aggagaaccc cagggctgcc tggctcccca tagccaaccc catccccaat
                                                                    1020
ttccagtgtg gcaccctgcc tgagaccggt cccaacgaga acctgacgga gcgcagcctg
caggacgcgc agcgcctctt cctgatgagc gaggccgtgc agccggtgac acccgagccc
                                                                    1080
                                                                    1140
tgtgtcaccc aggacagcgt gcgcttctca cacctcgtgg tggacctggt gcaggctaaa
                                                                    1200
gacacgetet accatgtact etacattgge accgagtegg geaccateet gaaggegetg
tccacggcga gccgcagcct ccacggctgc tacctggagg agctgcacgt gctgccccc
                                                                    1260
                                                                    1320
gggcgccgcg agcccctgcg cagcctgcgc atcctgcaca gcgcccgcgc gctcttcgtg
gggctgagag acggcgtcct gcgggtccca ctggagaggt gcgccgccta ccgcagccag
                                                                    1380
                                                                    1440
ggggcatgcc tgggggcccg ggacccgtac tgtggctggg acgggaagca gcaacgttgc
                                                                    1500
agcacactcg aggacagete caacatgage etetggacee agaacateae egeetgteet
gtgcggaatg tgacacggga tgggggcttc ggcccatggt caccatggca accatgtgag
                                                                    1560
cacttggatg gggacaactc aggetettge etgtgtegag etegateetg tgatteeect
                                                                    1620
                                                                    1680
cgaccccgct gtgggggcct tgactgcctg gggccagcca tccacatcgc caactgctcc
aggaatgggg cgtggacccc gtggtcatcg tgggcgctgt gcagcacgtc ctgtggcatc
                                                                    1740
```

```
ggcttccagg tccgccagcg aagttgcagc aaccctgctc cccgccacgg gggccgcatc
                                                                     1800
tgcgtgggca agagccggga ggaacggttc tgtaatgaga acacgccttg cccggtgccc
                                                                     1860
atcttctggg cttcctgggg ctcctggagc aagtgcagca gcaactgtgg agggggcatg
                                                                     1920
cagtcgcggc gtcgggcctg cgagaacggc aactcctgcc tgggctgcgg cgtggagttc
                                                                     1980
                                                                     2040
aagacgtgca accccgaggg ctgccccgaa gtgcggcgca acaccccctg gacgccgtgg
                                                                     2100
ctqccqtqa acqtgacqca gggcggggca cggcaggagc agcggttccg cttcacctgc
                                                                     2160
cgcgcgcccc ttgcagaccc gcacggcctg cagttcggca ggagaaggac cgagacgagg
                                                                     2220
acctgtcccg cggacggctc cggctcctgc gacaccgacg ccctggtgga ggacctcctg
                                                                     2280
cgcagcggga gcacctcccc gcacacggtg agcgggggct gggccgcctg gggcccgtgg
tcgtcctgct cccgggactg cgagctgggc ttccgcgtcc gcaagagaac gtgcactaac
                                                                     2340
                                                                     2400
ccqqaqcccc gcaacggggg cctgccctgc gtgggcgatg ctgccgagta ccaggactgc
                                                                     2460
aaccccagg cttgcccagt tcggggtgct tggtcctgct ggacctcatg gtctccatgc
tcagcttcct gtggtggggg tcactatcaa cgcacccgtt cctgcaccag ccccgcaccc
                                                                     2520
tccccaggtg aggacatctg tctcgggctg cacacggagg aggcactatg tgccacacag
                                                                     2580
gcctgcccag aaggctggtc gccctggtct gagtggagta agtgcactga cgacggagcc
                                                                     2640
                                                                     2700
cagagccgaa gccggcactg tgaggagctc ctcccagggt ccagcgcctg tgctggaaac
ageagecaga geogecett ceetacage gagatteeeg teateetgee ageeteeage
                                                                     2760
atggaggagg ccaccggctg tgcagggttc aatctcatcc acttggtggc cacgggcatc
                                                                     2820
tectgettet tgggetetgg geteetgace etageagtgt acetgtettg ecageactge
                                                                     2880
cagcgtcagt cccaggagtc cacactggtc catcctgcca cccccaacca tttgcactac
                                                                     2940
aagggcggag gcaccccgaa gaatgaaaag tacacaccca tggaattcaa gaccctgaac
                                                                     3000
aagaataact tgatccctga tgacagagcc aacttctacc cattgcagca gaccaatgtg
                                                                     3060
                                                                     3120
tacacgacta cttactaccc aagccccctg aacaaacaca gcttccggcc cgaggcctca
                                                                     3150
cctggacaac ggtgcttccc caacagctga
```

<210> 2 <211> 1049

<212> PRT

<213> homo sapiens

<400> 2

Met Thr Val Val Asn Pro Gln Asp Leu Gln Pro Trp Val Ser Asn Phe 10 Thr Tyr Pro Gly Ala Arg Asp Phe Ser Gln Leu Ala Leu Asp Pro Ser 2.5 20 Gly Asn Gln Leu Ile Val Gly Ala Arg Asn Tyr Leu Phe Arg Leu Ser 40 Leu Ala Asn Val Ser Leu Leu Gln Ala Thr Glu Trp Ala Ser Ser Glu 60 55 Asp Thr Arg Arg Ser Cys Gln Ser Lys Gly Lys Thr Glu Glu Glu Cys 75 80 70 Gln Asn Tyr Val Arg Val Leu Ile Val Ala Gly Arg Lys Val Phe Met 85 90 Cys Gly Thr Asn Ala Phe Ser Pro Met Cys Thr Ser Arg Gln Val Gly 110 105 100 Asn Leu Ser Arg Thr Ile Glu Lys Ile Asn Gly Val Ala Arg Cys Pro 125 120 Tyr Asp Pro Arg His Asn Ser Thr Ala Val Ile Ser Ser Gln Gly Glu 135 Leu Tyr Ala Ala Thr Val Ile Asp Phe Ser Gly Arg Asp Pro Ala Ile 155 150 Tyr Arg Ser Leu Gly Ser Gly Pro Pro Leu Arg Thr Ala Gln Tyr Asn 170 165 Ser Lys Trp Leu Asn Glu Pro Asn Phe Val Ala Ala Tyr Asp Ile Gly 185 190 Leu Phe Ala Tyr Phe Phe Leu Arg Glu Asn Ala Val Glu His Asp Cys

		195					200					205			
Gly	Arg 210	Thr	Val	Tyr	Ser	Arg 215	Val	Ala	Arg	Val	Cys 220	Lys	Asn	Asp	Val
Gly 225	Gly	Arg	Phe	Leu	Leu 230	Glu	Asp	Thr	Trp	Thr 235	Thr	Phe	Met	Lys	Ala 240
Arg	Leu	Asn	Cys	Ser 245	Arg	Pro	Gly	Glu	Val 250	Pro	Phe	Tyr	Tyr	Asn 255	Glu
Leu	Gln	Ser	Ala 260	Phe	His	Leu	Pro	Glu 265	Gln	Asp	Leu	I1e	Tyr 270	Gly	Val
Phe	Thr	Thr 275	Asn	Va1	Asn	Ser	Ile 280	Ala	Ala	Ser	Ala	Val 285	Cys	Ala	Phe
Asn	Leu 290	Ser	Ala	Ile	Ser	Gln 295	Ala	Phe	Asn	Gly	Pro 300	Phe	Arg	Tyr	Gln
305	Asn		_		310	_				315					320
	Gln			325					330					335	
Glu	Arg	Ser	Leu 340	Gln	Asp	Ala	Gln	Arg 345	Leu	Phe	Leu	Met	Ser 350	Glu	Ala
	Gln	355					360					365			
	Ser 370					375					380				
385	Val				390					395					400
	Thr			405					410					415	
	Leu		420	_	-	_		425					430		
	Ser	435					440					445			
	Pro 450					455					460				
465	Ala				470					475					480
	Thr			485					490					495	
	Ala		500					505					510		
	Ser	515					520					525			
	Cys 530		_			535					540				
545	Gly				550					555					560
	Asn			565					570					575	
	Cys		580					585					590		
	Pro	595					600					605			
	Phe 610					615					620				
625	Trp				630					635					640
Gln	Ser	Arg	Arg	Arg	Ala	Cys	Glu	Asn	G1y	Asn	Ser	Cys	Leu	Gly	Cys

				645					650					655	
Gly	Val	Glu	Phe 660	Lys	Thr	Cys	Asn	Pro 665	Glu	Gly	Cys	Pro	Glu 670	Val	Arg
Arg	Asn	Thr 675	Pro	Trp	Thr	Pro	Trp 680	Leu	Pro	Val	Asn	Val 685	Thr	Gln	Gly
G1y	Ala 690	Arg	Gln	Glu	Gln	Arg 695	Phe	Arg	Phe	Thr	Cys 700	Arg	Ala	Pro	Leu
Ala 705	Asp	Pro	His	Gly	Leu 710	Gln	Phe	Gly	Arg	Arg 715	Arg	Thr	Glu	Thr	Arg 720
Thr	Cys	Pro	Ala	Asp 725	G1y	Ser	Gly	Ser	Cys 730	Asp	Thr	Asp	Ala	Leu 735	Val
Glu	Asp	Leu	Leu 740	Arg	Ser	Gly	Ser	Thr 745	Ser	Pro	His	Thr	Val 750	Ser	Gly
Gly	Trp	Ala 755	Ala	Trp	Gly	Pro	Trp 760	Ser	Ser	Cys	Ser	Arg 765	Asp	Cys	Glu
	770					775					780			Pro	
Asn 785	Gly	Gly	Leu	Pro	Cys 790	Val	Gly	Asp	Ala	Ala 795	Glu	Tyr	Gln	Asp	800
				805					810					Thr 815	
Trp	Ser	Pro	Cys 820	Ser	Ala	Ser	Суз	Gly 825	Gly	Gly	His	Tyr	Gln 830	Arg	Thr
Arg	Ser	Cys 835	Thr	Ser	Pro	Ala	Pro 840	Ser	Pro	Gly	Glu	Asp 845	Ile	Cys	Leu
Gly	Leu 850	His	Thr	Glu	Glu	Ala 855	Leu	Суѕ	Ala	Thr	Gln 860	Ala	Суз	Pro	Glu
Gly 865	Trp	Ser	Pro	Trp	Ser 870	Glu	Trp	Ser	Lys	Суs 875	Thr	Asp	Asp	Gly	Ala 880
Gln	Ser	Arg	Ser	Arg 885	His	Cys	Glu	Glu	Leu 890	Leu	Pro	Gly	Ser	Ser 895	Ala
Суз	Ala	Gly	Asn 900	Ser	Ser	Gln	Ser	Arg 905	Pro	Cys	Pro	Tyr	Ser 910	Glu	Ile
Pro	Val	Ile 915	Leu	Pro	Ala	Ser	Ser 920	Met	Glu	Glu	Ala	Thr 925	Gly	Cys	Ala
Gly	Phe 930	Asn	Leu	Ile	His	Leu 935	Va1	Ala	Thr	Gly	Ile 940	Ser	Cys	Phe	Leu
Gly 945	Ser	Gly	Leu	Leu	Thr 950	Leu	Ala	Val	Tyr	Leu 955	Ser	Cys	Gln	His	Cys 960
Gln	Arg	Gln	Ser	Gln 965	Glu	Ser	Thr	Leu	Val 970	His	Pro	Ala	Thr	Pro 975	Asn
His	Leu	His	Tyr 980	Lys	Gly	G1y	Gly	Thr 985	Pro	Lys	Asn	Glu	Lys 990	Tyr	Thr
Pro	Met	Glu 995	Phe	Lys	Thr	Leu	Asn 100		Asn	Asn	Leu	Ile 100		Asp	Asp
Arg	Ala 101		Phe	Tyr	Pro	Leu 101!		Gln	Thr	Asn	Val 102		Thr	Thr	Thr
Tyr 1025	_	Pro	Ser	Pro	Leu 103		Lys	His	Ser	Phe 103		Pro	Glu	Ala	Ser 1040
Pro	Gly	Gln	Arg	Cys 104		Pro	Asn	Ser							

<210> 3 <211> 3282

<212> DNA

<213> homo sapiens

<400> 3						60
	caggeeecet					60 120
	ccagctccca					120
	agcaccccac					180
	gagcccggga					240
	ccaggaacta	_				300
	gggcctccag					360
	gtcagaacta					420
	atgccttttc					480
	agatcaatgg					540
gctgtcatct	cctcccaggg	ggagctctat	gcagccacgg	tcatcgactt	ctcaggtcgg	600
	tctaccgcag					660
tccaagtggc	ttaatgagcc	aaacttcgtg	gcagcctatg	atattgggct	gtttgcatac	720
ttcttcctgc	gggagaacgc	agtggagcac	gactgtggac	gcaccgtgta	ctctcgcgtg	780
	gcaagaatga					840
ttcatgaagg	cccggctcaa	ctgctcccgc	ccgggcgagg	tccccttcta	ctataacgag	900
ctgcagagtg	ccttccactt	gccrgagcag	gacctcatct	atggagtttt	cacaaccaac	960
gtaaacagca	tygcggcttc	tgctgtctgc	gccttcaacc	tcagtgctat	ctcccaggct	1020
ttcaatggcc	catttcgcta	ccaggagaac	cccagggctg	cctggctccc	catagccaac	1080
cccatcccca	atttccagtg	tggcaccctg	cctgagaccg	gtcccaacga	gaacctgacg	1140
gagcgcagcc	tgcaggacgc	gcagcgcctc	ttcctgatga	gcgaggccgt	gcagccggtg	1200
acacccgagc	cctgtgtcac	ccaggacagc	gtgcgcttct	cacacctcgt	ggtggacctg	1260
gtgcaggcta	aagacacgct	ctaccatgta	ctctacattg	gcaccgagtc	gggcaccatc	1320
ctgaaggcgc	tgtccacggc	gagccgcagc	ctccacggct	gctacctgga	ggagctgcac	1380
gtgctgcccc	ccgggcgccg	cgagcccctg	cgcagcctgc	gcatcctgca	cagcgcccgc	1440
gcgctcttcg	tggggctgag	agacggcgtc	ctgcgggtcc	cactggagag	gtgcgccgcc	1500
	agggggcatg					1560
cagcaacgtt	gcagcacact	cgaggacagc	tccaacatga	gcctctggac	ccagaacatc	1620
	ctgtgcggaa					1680
caaccatgtg	agcacttgga	tggggacaac	tcaggctctt	gcctgtgtcg	agctcgatcc	1740
tgtgattccc	ctcgaccccg	ctgtgggggc	cttgactgcc	tggggccagc	catccacatc	1800
gccaactgct	ccaggaatgg	ggcgtggacc	ccgtggtcat	cgtgggcgct	gtgcagcacg	1860
tcctgtggca	tcggcttcca	ggtccgccag	cgaagttgca	gcaaccctgc	tccccgccac	1920
gggggccgca	tctgcgtggg	caagagccgg	gaggaacggt	tctgtaatga	gaacacgcct	1980
tgcccggtgc	ccatcttctg	ggcttcctgg	ggctcctgga	gcaagtgcag	cagcaactgt	2040
ggagggggca	tgcagtcgcg	gcgtcgggcc	tgcgagaacg	gcaactcctg	cctgggctgc	2100
ggcgtggagt	tcaagacgtg	caaccccgag	ggctgccccg	aagtgcggcg	caacaccccc	2160
	ggctgcccgt					2220
	gccgcgcgcc					2280
	ggacctgtcc					2340
	tgcgcagcgg					2400
	ggtcgtcctg					2460
	acccggagcc					2520
	gcaaccccca					2580
	gctcagcttc					2640
	cctccccagg					2700
	aggcctgccc					2760
	cccagagccg					2820
	acagcagcca					2880
	gcatggagga					2940
	tctcctgctt					3000
	gccagcgtca					3060
	acaagggcgg					3120
	acaagaataa					3180
	tgtacacgac					3240
	cacctggaca				. 5 55	3282
2 23 4-	JJ	33 3	5 5			

<210> 4 <211> 1093

```
<212> PRT
<213> homo sapiens
<400> 4
Met Val Leu Ala Gly Pro Leu Ala Val Ser Leu Leu Pro Ser Leu
                                  1.0
Thr Leu Leu Val Ser His Leu Ser Ser Gln Asp Val Ser Ser Glu
                              25
Pro Ser Ser Glu Gln Gln Leu Cys Ala Leu Ser Lys His Pro Thr Val
                          40
Ala Phe Glu Asp Leu Gln Pro Trp Val Ser Asn Phe Thr Tyr Pro Gly
Ala Arg Asp Phe Ser Gln Leu Ala Leu Asp Pro Ser Gly Asn Gln Leu
                  70
                                     75
Ile Val Gly Ala Arg Asn Tyr Leu Phe Arg Leu Ser Leu Ala Asn Val
                                 90
Ser Leu Leu Gln Ala Thr Glu Trp Ala Ser Ser Glu Asp Thr Arg Arg
                             105
                                                110
          100
Ser Cys Gln Ser Lys Gly Lys Thr Glu Glu Glu Cys Gln Asn Tyr Val
                         120
                                             125
Arg Val Leu Ile Val Ala Gly Arg Lys Val Phe Met Cys Gly Thr Asn
                      135
Ala Phe Ser Pro Met Cys Thr Ser Arg Gln Val Gly Asn Leu Ser Arg
                   150
                                      155
Thr Ile Glu Lys Ile Asn Gly Val Ala Arg Cys Pro Tyr Asp Pro Arg
               165
                                  170
His Asn Ser Thr Ala Val Ile Ser Ser Gln Gly Glu Leu Tyr Ala Ala
                              185
  180
                                                 190
Thr Val Ile Asp Phe Ser Gly Arg Asp Pro Ala Ile Tyr Arg Ser Leu
                         200
Gly Ser Gly Pro Pro Leu Arg Thr Ala Gln Tyr Asn Ser Lys Trp Leu
                      215
                                         220
Asn Glu Pro Asn Phe Val Ala Ala Tyr Asp Ile Gly Leu Phe Ala Tyr
     230
                                      235
Phe Phe Leu Arg Glu Asn Ala Val Glu His Asp Cys Gly Arg Thr Val
                                 250
              245
Tyr Ser Arg Val Ala Arg Val Cys Lys Asn Asp Val Gly Gly Arg Phe
                             265
Leu Leu Glu Asp Thr Trp Thr Thr Phe Met Lys Ala Arg Leu Asn Cys
                           280
Ser Arg Pro Gly Glu Val Pro Phe Tyr Tyr Asn Glu Leu Gln Ser Ala
                       295
Phe His Leu Pro Glu Gln Asp Leu Ile Tyr Gly Val Phe Thr Thr Asn
                   310
                                      315
Val Asn Ser Ile Ala Ala Ser Ala Val Cys Ala Phe Asn Leu Ser Ala
                                  330
               325
Ile Ser Gln Ala Phe Asn Gly Pro Phe Arg Tyr Gln Glu Asn Pro Arg
                             345
Ala Ala Trp Leu Pro Ile Ala Asn Pro Ile Pro Asn Phe Gln Cys Gly
                          360
                                              365
Thr Leu Pro Glu Thr Gly Pro Asn Glu Asn Leu Thr Glu Arg Ser Leu
                       375
                                         380
Gln Asp Ala Gln Arg Leu Phe Leu Met Ser Glu Ala Val Gln Pro Val
                   390
                                      395
```

Thr Pro Glu Pro Cys Val Thr Gln Asp Ser Val Arg Phe Ser His Leu 405 410 Val Val Asp Leu Val Gln Ala Lys Asp Thr Leu Tyr His Val Leu Tyr 425 Ile Gly Thr Glu Ser Gly Thr Ile Leu Lys Ala Leu Ser Thr Ala Ser 440 Arg Ser Leu His Gly Cys Tyr Leu Glu Glu Leu His Val Leu Pro Pro 455 Gly Arg Arg Glu Pro Leu Arg Ser Leu Arg Ile Leu His Ser Ala Arg 475 470 Ala Leu Phe Val Gly Leu Arg Asp Gly Val Leu Arg Val Pro Leu Glu Arg Cys Ala Ala Tyr Arg Ser Gln Gly Ala Cys Leu Gly Ala Arg Asp 505 Pro Tyr Cys Gly Trp Asp Gly Lys Gln Gln Arg Cys Ser Thr Leu Glu 520 Asp Ser Ser Asn Met Ser Leu Trp Thr Gln Asn Ile Thr Ala Cys Pro 535 540 Val Arg Asn Val Thr Arg Asp Gly Gly Phe Gly Pro Trp Ser Pro Trp 550 555 Gln Pro Cys Glu His Leu Asp Gly Asp Asn Ser Gly Ser Cys Leu Cys 570 Arg Ala Arg Ser Cys Asp Ser Pro Arg Pro Arg Cys Gly Gly Leu Asp 580 585 Cys Leu Gly Pro Ala Ile His Ile Ala Asn Cys Ser Arg Asn Gly Ala 600 Trp Thr Pro Trp Ser Ser Trp Ala Leu Cys Ser Thr Ser Cys Gly Ile 615 620 Gly Phe Gln Val Arg Gln Arg Ser Cys Ser Asn Pro Ala Pro Arg His 630 635 Gly Gly Arg Ile Cys Val Gly Lys Ser Arg Glu Glu Arg Phe Cys Asn 645 650 Glu Asn Thr Pro Cys Pro Val Pro Ile Phe Trp Ala Ser Trp Gly Ser 665 660 Trp Ser Lys Cys Ser Ser Asn Cys Gly Gly Met Gln Ser Arg Arg 680 Arg Ala Cys Glu Asn Gly Asn Ser Cys Leu Gly Cys Gly Val Glu Phe 695 Lys Thr Cys Asn Pro Glu Gly Cys Pro Glu Val Arg Arg Asn Thr Pro 710 715 Trp Thr Pro Trp Leu Pro Val Asn Val Thr Gln Gly Gly Ala Arg Gln 730 Glu Gln Arg Phe Arg Phe Thr Cys Arg Ala Pro Leu Ala Asp Pro His 745 Gly Leu Gln Phe Gly Arg Arg Thr Glu Thr Arg Thr Cys Pro Ala 760 Asp Gly Ser Gly Ser Cys Asp Thr Asp Ala Leu Val Glu Asp Leu Leu 775 Arg Ser Gly Ser Thr Ser Pro His Thr Val Ser Gly Gly Trp Ala Ala 790 795 Trp Gly Pro Trp Ser Ser Cys Ser Arg Asp Cys Glu Leu Gly Phe Arg 810 Val Arg Lys Arg Thr Cys Thr Asn Pro Glu Pro Arg Asn Gly Gly Leu 825 820 Pro Cys Val Gly Asp Ala Ala Glu Tyr Gln Asp Cys Asn Pro Gln Ala 840

```
Cys Pro Val Arg Gly Ala Trp Ser Cys Trp Thr Ser Trp Ser Pro Cys
                       855
Ser Ala Ser Cys Gly Gly His Tyr Gln Arg Thr Arg Ser Cys Thr
                                      875
                   870
Ser Pro Ala Pro Ser Pro Gly Glu Asp Ile Cys Leu Gly Leu His Thr
                                   890
Glu Glu Ala Leu Cys Ala Thr Gln Ala Cys Pro Glu Gly Trp Ser Pro
                               905
Trp Ser Glu Trp Ser Lys Cys Thr Asp Asp Gly Ala Gln Ser Arg Ser
                           920
Arg His Cys Glu Glu Leu Leu Pro Gly Ser Ser Ala Cys Ala Gly Asn
                       935
Ser Ser Gln Ser Arg Pro Cys Pro Tyr Ser Glu Ile Pro Val Ile Leu
                   950
                                       955
Pro Ala Ser Ser Met Glu Glu Ala Thr Gly Cys Ala Gly Phe Asn Leu
                                   970
               965
Ile His Leu Val Ala Thr Gly Ile Ser Cys Phe Leu Gly Ser Gly Leu
           980
                              985
Leu Thr Leu Ala Val Tyr Leu Ser Cys Gln His Cys Gln Arg Gln Ser
                           1000
                                              1005
Gln Glu Ser Thr Leu Val His Pro Ala Thr Pro Asn His Leu His Tyr
                      1015
                                          1020
Lys Gly Gly Gly Thr Pro Lys Asn Glu Lys Tyr Thr Pro Met Glu Phe
                   1030
                                      1035
Lys Thr Leu Asn Lys Asn Asn Leu Ile Pro Asp Asp Arg Ala Asn Phe
               1045
                                   1050
Tyr Pro Leu Gln Gln Thr Asn Val Tyr Thr Thr Tyr Tyr Pro Ser
                               1065
                                                  1070
Pro Leu Asn Lys His Ser Phe Arg Pro Glu Ala Ser Pro Gly Gln Arg
                           1080
       1075
                                               1085
Cys Phe Pro Asn Ser
    1090
<210> 5
<211> 3105
<212> DNA
<213> homo sapiens
<400> 5
                                                                     60
atgacggtgg ttaatccaca agacctgcag ccgtgggtct ctaacttcac ctaccctgga
                                                                    120
geeegggatt teteceaget ggetttggae eeeteegggr accageteat egtgggagee
                                                                    180
aggaactacc tetteagact cageettgee aatgtetete ttetteagge cacagagtgg
                                                                    240
300
cagaactacg tgcgagtcct gatcgtcgcc ggccggaagg tgttcatgtg tggaaccaat
                                                                    360
gccttttccc ccatgtgcac cagcagacag gtggggaacc tcagccggac tattgagaag
                                                                    420
atcaatggtg tggcccgctg cccctatgac ccacgccaca actccacagc tgtcatctcc
teccaggggg agetetatge agecaeggte ategaettet caggteggga ccctgccate
                                                                    480
                                                                    540
taccgcagcc tgggcagtgg gccaccgctt cgcactgccc aatataactc caagtggctt
                                                                    600
aatgagccaa acttcgtggc agcctatgat attgggctgt ttgcatactt cttcctgcgg
                                                                    660
gagaacgcag tggagcacga ctgtggacgc accgtgtact ctcgcgtggc ccgcgtgtgc
                                                                    720
aagaatgacg tggggggcg attectgctg gaggacacat ggaccacatt catgaaggcc
eggeteaact geteeegeee gggegaggte eeettetaet ataacgaget geagagtgee
                                                                    780
ttccacttgc crgagcagga cctcatctat ggagttttca caaccaacgt aaacagcaty
                                                                    840
                                                                    900
geggettetg etgtetgege etteaacete agtgetatet eccaggettt caatggeeca
tttcgctacc aggagaaccc cagggctgcc tggctcccca tagccaaccc catccccaat
                                                                    960
ttccagtgtg gcaccctgcc tgagaccggt cccaacgaga acctgacgga gcgcagcctg
                                                                   1020
```

```
caggacgcgc agcgcctctt cctgatgagc gaggccgtgc agccggtgac acccgagccc
                                                                     1080
                                                                     1140
tgtgtcaccc aggacagcgt gcgcttctca cacctcgtgg tggacctggt gcaggctaaa
gacacgctct accatgtact ctacattggc accgagtcgg gcaccatcct gaaggcgctg
                                                                     1200
tocacggega geogrageet ceaeggetge tacetggagg agetgeaegt getgeeeee
                                                                     1260
gggcgccgcg agcccctgcg cagcctgcgc atcctgcaca gcgcccgcgc gctcttcgtg
                                                                     1320
gggctgagag acggcgtcct gcgggtccca ctggagaggt gcgccgccta ccgcagccag
                                                                     1380
ggggcatgcc tgggggcccg ggacccgtac tgtggctggg acgggaagca gcaacgttgc
                                                                     1440
agcacactcg aggacagete caacatgage etetggacee agaacateae egeetgteet
                                                                     1500
                                                                     1560
gtgcggaatg tgacacggga tgggggcttc ggcccatggt caccatggca accatgtgag
                                                                     1620
cacttggatg gggacaactc aggctettge etgtgtegag etegateetg tgatteeeet
                                                                     1680
cgaccccgct gtgggggcct tgactgcctg gggccagcca tccacatcgc caactgctcc
                                                                     1740
aggaatgggg cgtggacccc gtggtcatcg tgggcgctgt gcagcacgtc ctgtggcatc
                                                                     1800
ggcttccagg tccgccagcg aagttgcagc aaccetgete cccgccacgg gggccgcate
tgcgtgggca agagccggga ggaacggttc tgtaatgaga acacgccttg cccggtgccc
                                                                     1860
                                                                     1920
atcttetggg etteetgggg eteetggage aagtgeagea geaactgtgg agggggeatg
cagtcgcggc gtcgggcctg cgagaacggc aactcctgcc tgggctgcgg cgtggagttc
                                                                     1980
aagacgtgca accccgaggg ctgccccgaa gtgcggcgca acaccccctg gacgccgtgg
                                                                     2040
ctgcccgtga acgtgacgca gggcggggca cggcaggagc agcggttccg cttcacctgc
                                                                      2100
                                                                     2160
cgcgcgcccc ttgcagaccc gcacggcctg cagttcggca ggagaaggac cgagacgagg
                                                                     2220
acctgtcccg cggacggctc cggctcctgc gacaccgacg ccctggtgga ggacctcctg
                                                                     2280
cgcagcggga gcacctcccc gcacacggtg agcgggggct gggccgcctg gggcccgtgg
tegteetget eeegggactg egagetggge tteegegtee geaagagaae gtgeactaae
                                                                     2340
ccggagcccc gcaacggggg cctgccctgc gtgggcgatg ctgccgagta ccaggactgc
                                                                     2400
aacccccagg cttgcccagt tcggggtgct tggtcctgct ggacctcatg gtctccatgc
                                                                     2460
                                                                     2520
tragetteet gtggtggggg teactateaa egeaceegtt cetgeaceag cecegeacee
                                                                     2580
tccccaggtg aggacatctg tctcgggctg cacacggagg aggcactatg tgccacacag
gcctgcccag aaggctggtc gccctggtct gagtggagta agtgcactga cgacggagcc
                                                                     2640
                                                                     2700
cagageegaa geeggeactg tgaggagete etceeagggt ecagegeetg tgetggaaac
                                                                     2760
agcagecaga geogecectg eccetacage gagatteeeg ggtteaatet catecaettg
                                                                     2820
gtggccacgg gcatctcctg cttcttgggc tctgggctcc tgaccctagc agtgtacctg
tottgccage actgccageg teagteccag gagtecaeae tggtccatee tgccaeeeee
                                                                     2880
                                                                     2940
aaccatttgc actacaaggg cggaggcacc ccgaagaatg aaaagtacac acccatggaa
                                                                     3000
ttcaagaccc tgaacaagaa taacttgatc cctgatgaca gagccaactt ctacccattg
                                                                     3060
cagcagacca atgtgtacac gactacttac tacccaagcc ccctgaacaa acacagcttc
                                                                     3105
cggcccgagg cctcacctgg acaacggtgc ttccccaaca gctga
```

```
<210> 6
```

## <400> 6

```
Met Thr Val Val Asn Pro Gln Asp Leu Gln Pro Trp Val Ser Asn Phe
1
                 5
                                     10
Thr Tyr Pro Gly Ala Arg Asp Phe Ser Gln Leu Ala Leu Asp Pro Ser
                                 25
Gly Asn Gln Leu Ile Val Gly Ala Arg Asn Tyr Leu Phe Arg Leu Ser
                             40
Leu Ala Asn Val Ser Leu Leu Gln Ala Thr Glu Trp Ala Ser Ser Glu
                        55
                                             60
Asp Thr Arg Arg Ser Cys Gln Ser Lys Gly Lys Thr Glu Glu Glu Cys
                    70
                                         75
Gln Asn Tyr Val Arg Val Leu Ile Val Ala Gly Arg Lys Val Phe Met
                                     90
Cys Gly Thr Asn Ala Phe Ser Pro Met Cys Thr Ser Arg Gln Val Gly
            100
                                105
```

<sup>&</sup>lt;211> 1034

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> homo sapiens

```
Asn Leu Ser Arg Thr Ile Glu Lys Ile Asn Gly Val Ala Arg Cys Pro
                           120
Tyr Asp Pro Arg His Asn Ser Thr Ala Val Ile Ser Ser Gln Gly Glu
                      135
                                           140
Leu Tyr Ala Ala Thr Val Ile Asp Phe Ser Gly Arg Asp Pro Ala Ile
                   150
                                       155
Tyr Arg Ser Leu Gly Ser Gly Pro Pro Leu Arg Thr Ala Gln Tyr Asn
                                   170
               165
Ser Lys Trp Leu Asn Glu Pro Asn Phe Val Ala Ala Tyr Asp Ile Gly
                               185
           180
Leu Phe Ala Tyr Phe Phe Leu Arg Glu Asn Ala Val Glu His Asp Cys
                            200
Gly Arg Thr Val Tyr Ser Arg Val Ala Arg Val Cys Lys Asn Asp Val
                      215
Gly Gly Arg Phe Leu Leu Glu Asp Thr Trp Thr Thr Phe Met Lys Ala
                  230
                                       235
Arg Leu Asn Cys Ser Arg Pro Gly Glu Val Pro Phe Tyr Tyr Asn Glu
                                    250
               245
Leu Gln Ser Ala Phe His Leu Pro Glu Gln Asp Leu Ile Tyr Gly Val
           260
                               265
Phe Thr Thr Asn Val Asn Ser Ile Ala Ala Ser Ala Val Cys Ala Phe
                                               285
                           280
Asn Leu Ser Ala Ile Ser Gln Ala Phe Asn Gly Pro Phe Arg Tyr Gln
                                           300
                        295
Glu Asn Pro Arg Ala Ala Trp Leu Pro Ile Ala Asn Pro Ile Pro Asn
                    310
                                        315
Phe Gln Cys Gly Thr Leu Pro Glu Thr Gly Pro Asn Glu Asn Leu Thr
                                    330
               325
Glu Arg Ser Leu Gln Asp Ala Gln Arg Leu Phe Leu Met Ser Glu Ala
                               345
Val Gln Pro Val Thr Pro Glu Pro Cys Val Thr Gln Asp Ser Val Arg
                           360
       355
Phe Ser His Leu Val Val Asp Leu Val Gln Ala Lys Asp Thr Leu Tyr
                       375
                                            380
His Val Leu Tyr Ile Gly Thr Glu Ser Gly Thr Ile Leu Lys Ala Leu
                                       395
                    390
Ser Thr Ala Ser Arg Ser Leu His Gly Cys Tyr Leu Glu Glu Leu His
                405
Val Leu Pro Pro Gly Arg Arg Glu Pro Leu Arg Ser Leu Arg Ile Leu
                                425
His Ser Ala Arg Ala Leu Phe Val Gly Leu Arg Asp Gly Val Leu Arg
                            440
Val Pro Leu Glu Arg Cys Ala Ala Tyr Arg Ser Gln Gly Ala Cys Leu
                       455
Gly Ala Arg Asp Pro Tyr Cys Gly Trp Asp Gly Lys Gln Gln Arg Cys
                   470
                                       475
Ser Thr Leu Glu Asp Ser Ser Asn Met Ser Leu Trp Thr Gln Asn Ile
                485
                                    490
Thr Ala Cys Pro Val Arg Asn Val Thr Arg Asp Gly Phe Gly Pro
            500
                                505
Trp Ser Pro Trp Gln Pro Cys Glu His Leu Asp Gly Asp Asn Ser Gly
                            520
Ser Cys Leu Cys Arg Ala Arg Ser Cys Asp Ser Pro Arg Pro Arg Cys
                        535
                                           540
Gly Gly Leu Asp Cys Leu Gly Pro Ala Ile His Ile Ala Asn Cys Ser
                                        555
545
                    550
```

```
Arg Asn Gly Ala Trp Thr Pro Trp Ser Ser Trp Ala Leu Cys Ser Thr
               565
                                   570
Ser Cys Gly Ile Gly Phe Gln Val Arg Gln Arg Ser Cys Ser Asn Pro
                               585
           580
Ala Pro Arg His Gly Gly Arg Ile Cys Val Gly Lys Ser Arg Glu Glu
                                              605
                           600
Arg Phe Cys Asn Glu Asn Thr Pro Cys Pro Val Pro Ile Phe Trp Ala
                                           620
                      615
Ser Trp Gly Ser Trp Ser Lys Cys Ser Ser Asn Cys Gly Gly Met
                                      635
                   630
Gln Ser Arg Arg Ala Cys Glu Asn Gly Asn Ser Cys Leu Gly Cys
                                   650
Gly Val Glu Phe Lys Thr Cys Asn Pro Glu Gly Cys Pro Glu Val Arg
                               665
Arg Asn Thr Pro Trp Thr Pro Trp Leu Pro Val Asn Val Thr Gln Gly
                           680
Gly Ala Arg Gln Glu Gln Arg Phe Arg Phe Thr Cys Arg Ala Pro Leu
                       695
Ala Asp Pro His Gly Leu Gln Phe Gly Arg Arg Arg Thr Glu Thr Arg
                                       715
                    710
Thr Cys Pro Ala Asp Gly Ser Gly Ser Cys Asp Thr Asp Ala Leu Val
                                   730
                725
Glu Asp Leu Leu Arg Ser Gly Ser Thr Ser Pro His Thr Val Ser Gly
                               745
            740
Gly Trp Ala Ala Trp Gly Pro Trp Ser Ser Cys Ser Arg Asp Cys Glu
                            760
        755
Leu Gly Phe Arg Val Arg Lys Arg Thr Cys Thr Asn Pro Glu Pro Arg
                                           780
                        775
Asn Gly Gly Leu Pro Cys Val Gly Asp Ala Ala Glu Tyr Gln Asp Cys
                                       795
                   790
Asn Pro Gln Ala Cys Pro Val Arg Gly Ala Trp Ser Cys Trp Thr Ser
                                   810
               805
Trp Ser Pro Cys Ser Ala Ser Cys Gly Gly Gly His Tyr Gln Arg Thr
                              825
           820
 Arg Ser Cys Thr Ser Pro Ala Pro Ser Pro Gly Glu Asp Ile Cys Leu
                           840
 Gly Leu His Thr Glu Glu Ala Leu Cys Ala Thr Gln Ala Cys Pro Glu
                              860
 Gly Trp Ser Pro Trp Ser Glu Trp Ser Lys Cys Thr Asp Asp Gly Ala
                                        875
                    870
Gln Ser Arg Ser Arg His Cys Glu Glu Leu Leu Pro Gly Ser Ser Ala
                                    890
                885
 Cys Ala Gly Asn Ser Ser Gln Ser Arg Pro Cys Pro Tyr Ser Glu Ile
                               905
 Pro Gly Phe Asn Leu Ile His Leu Val Ala Thr Gly Ile Ser Cys Phe
                            920
 Leu Gly Ser Gly Leu Leu Thr Leu Ala Val Tyr Leu Ser Cys Gln His
                                           940
                       935
 Cys Gln Arg Gln Ser Gln Glu Ser Thr Leu Val His Pro Ala Thr Pro
                                        955
                    950
 Asn His Leu His Tyr Lys Gly Gly Gly Thr Pro Lys Asn Glu Lys Tyr
                                    970
                 965
 Thr Pro Met Glu Phe Lys Thr Leu Asn Lys Asn Asn Leu Ile Pro Asp
                                985
 Asp Arg Ala Asn Phe Tyr Pro Leu Gln Gln Thr Asn Val Tyr Thr Thr
                                                1005
                             1000
```

```
Thr Tyr Tyr Pro Ser Pro Leu Asn Lys His Ser Phe Arg Pro Glu Ala
                                            1020
                        1015
Ser Pro Gly Gln Arg Cys Phe Pro Asn Ser
1025
                    1030
<210> 7
<211> 3237
<212> DNA
<213> homo sapiens
<400> 7
                                                                        60
atggtgcttg caggcccct ggctgtctcg ctgttgctgc ccagcctcac actgctggtg
                                                                       120
tcccacctct ccagctccca ggatgtctcc agtgagccca gcagtgagca gcagctgtgc
                                                                       180
qcccttagca agcaccccac cgtggccttt gaagacctgc agccgtgggt ctctaacttc
acctaccctg gagcccggga tttctcccag ctggctttgg acccctccgg graccagctc
                                                                       240
                                                                       300
atcqtqqqaq ccaqqaacta cctcttcaga ctcagccttg ccaatgtctc tcttcttcag
                                                                       360
gccacagagt gggcctccag tgaggacacg cgccgctcct gccaaagcaa agggaagact
gaggaggagt gtcagaacta cgtgcgagtc ctgatcgtcg ccggccggaa ggtgttcatg
                                                                       420
                                                                       480
tqtqqaacca atgccttttc ccccatgtgc accagcagac aggtggggaa cctcagccgg
                                                                       540
actattgaga agatcaatgg tgtggcccgc tgcccctatg acccacgcca caactccaca
gctgtcatct cctcccaggg ggagctctat gcagccacgg tcatcgactt ctcaggtcgg
                                                                       600
gaccetgcca tetacegcag cetgggcagt gggccacege ttegcactge ccaatataac
                                                                       660
tccaagtggc ttaatgagcc aaacttcgtg gcagcctatg atattgggct gtttgcatac
                                                                       720
                                                                       780
ttetteetge gggagaacge agtggageae gaetgtggae geaeegtgta etetegegtg
gcccgcgtgt gcaagaatga cgtggggggc cgattcctgc tggaggacac atggaccaca
                                                                       840
ttcatgaagg cccggctcaa ctgctcccgc ccgggcgagg tccccttcta ctataacgag
                                                                       900
                                                                       960
ctgcagagtg ccttccactt gccrgagcag gacctcatct atggagtttt cacaaccaac
                                                                      1020
gtaaacagca tygcggcttc tgctgtctgc gccttcaacc tcagtgctat ctcccaggct
ttcaatggcc catttcgcta ccaggagaac cccagggctg cctggctccc catagccaac
                                                                      1080
                                                                      1140
cccatccca atttccaqtq tqqcaccctg cctgagaccg gtcccaacga gaacctgacg
                                                                      1200
gagcgcagcc tgcaggacgc gcagcgcctc ttcctgatga gcgaggccgt gcagccggtg
                                                                      1260
acacccgage cetgtgteac ceaggacage gtgegettet caeacctegt ggtggacetg
gtgcaggcta aagacacgct ctaccatgta ctctacattg gcaccgagtc gggcaccatc
                                                                      1320
                                                                      1380
ctgaaggcgc tgtccacggc gagccgcagc ctccacggct gctacctgga ggagctgcac
                                                                      1440
gtgctgcccc ccgggcgccg cgagcccctg cgcagcctgc gcatcctgca cagcgcccgc
                                                                      1500
gcgctcttcg tggggctgag agacggcgtc ctgcgggtcc cactggagag gtgcgccgcc
taccgcagcc agggggcatg cctgggggcc cgggacccgt actgtggctg ggacgggaag
                                                                      1560
cagcaacgtt gcagcacact cgaggacagc tccaacatga gcctctggac ccagaacatc
                                                                      1620
accgcctgtc ctgtgcggaa tgtgacacgg gatgggggct tcggcccatg gtcaccatgg
                                                                      1680
caaccatgtg agcacttgga tggggacaac tcaggetett geetgtgteg agetegatee
                                                                      1740
tgtgattccc ctcgaccccg ctgtgggggc cttgactgcc tgggggccagc catccacatc
                                                                      1800
gccaactgct ccaggaatgg ggcgtggacc ccgtggtcat cgtgggcgct gtgcagcacg
                                                                      1860
tcctgtggca tcggcttcca ggtccgccag cgaagttgca gcaaccctgc tccccgccac
                                                                      1920
qqqqqccqca tctqcqtqqq caaqaqccqq qagqaacqqt tctqtaatqa qaacacqcct
                                                                      1980
tgcccggtgc ccatcttctg ggcttcctgg ggctcctgga gcaagtgcag cagcaactgt
                                                                      2040
                                                                      2100
ggagggggca tgcagtcgcg gcgtcgggcc tgcgagaacg gcaactcctg cctgggctgc
                                                                      2160
ggcgtggagt tcaagacgtg caaccccgag ggctgccccg aagtgcggcg caacaccccc
tggacgccgt ggctgcccgt gaacgtgacg cagggcgggg cacggcagga gcagcggttc
                                                                      2220
                                                                      2280
cgcttcacct qccqcqccc ccttgcagac ccgcacggcc tgcagttcgg caggagaagg
                                                                      2340
accgagacga ggacctgtcc cgcggacggc tccggctcct gcgacaccga cgccctggtg
gaggacetee tgegeagegg gageacetee cegeacaegg tgageggggg etgggeegee
                                                                      2400
tggggcccgt ggtcgtcctg ctcccgggac tgcgagctgg gcttccgcgt ccgcaagaga
                                                                       2460
                                                                      2520
acgtgcacta acccggagec ccgcaacggg ggcctgccct gcgtgggcga tgctgccgag
taccaggact gcaaccccca ggcttgccca gttcggggtg cttggtcctg ctggacctca
                                                                      2580
tggtctccat gctcagcttc ctgtggtggg ggtcactatc aacgcacccg ttcctgcacc
                                                                      2640
```

agccccgcac cctccccagg tgaggacatc tgtctcgggc tgcacacgga ggaggcacta

```
tgtgccacac aggcctgccc agaaggctgg tcgccctggt ctgagtggag taagtgcact
gacgacggag cccagagccg aagccggcac tgtgaggagc tcctcccagg gtccagcgcc
tgtgctggaa acagcagcca gagccgcccc tgcccctaca gcgagattcc cgggttcaat
ctcatccact tggtggccac gggcatctcc tgcttcttgg gctctgggct cctgacccta
gcagtgtacc tgtcttgcca gcactgccag cgtcagtccc aggagtccac actggtccat
cctgccaccc ccaaccattt gcactacaag ggcggaggca ccccgaagaa tgaaaagtac
acacccatgg aattcaagac cctgaacaag aataacttga tccctgatga cagagccaac
ttctacccat tgcagcagac caatgtgtac acgactactt actacccaag ccccctgaac
aaacacagct tccggcccga ggcctcacct ggacaacggt gcttccccaa cagctga
<210> 8
<211> 1078
<212> PRT
<213> homo sapiens
<400> 8
Met Val Leu Ala Gly Pro Leu Ala Val Ser Leu Leu Leu Pro Ser Leu
                                    10
                5
Thr Leu Leu Val Ser His Leu Ser Ser Ser Gln Asp Val Ser Ser Glu
                                25
Pro Ser Ser Glu Gln Gln Leu Cys Ala Leu Ser Lys His Pro Thr Val
                            40
Ala Phe Glu Asp Leu Gln Pro Trp Val Ser Asn Phe Thr Tyr Pro Gly
                        55
                                            60
Ala Arg Asp Phe Ser Gln Leu Ala Leu Asp Pro Ser Gly Asn Gln Leu
                                        75
Ile Val Gly Ala Arg Asn Tyr Leu Phe Arg Leu Ser Leu Ala Asn Val
                                    90
                85
Ser Leu Leu Gln Ala Thr Glu Trp Ala Ser Ser Glu Asp Thr Arg Arg
                                105
Ser Cys Gln Ser Lys Gly Lys Thr Glu Glu Glu Cys Gln Asn Tyr Val
                            120
Arg Val Leu Ile Val Ala Gly Arg Lys Val Phe Met Cys Gly Thr Asn
                                            140
                        135
Ala Phe Ser Pro Met Cys Thr Ser Arg Gln Val Gly Asn Leu Ser Arg
                                       155
                    150
Thr Ile Glu Lys Ile Asn Gly Val Ala Arg Cys Pro Tyr Asp Pro Arg
                165
His Asn Ser Thr Ala Val Ile Ser Ser Gln Gly Glu Leu Tyr Ala Ala
                                 185
            180
Thr Val Ile Asp Phe Ser Gly Arg Asp Pro Ala Ile Tyr Arg Ser Leu
                            200
Gly Ser Gly Pro Pro Leu Arg Thr Ala Gln Tyr Asn Ser Lys Trp Leu
                                             220
                        215
Asn Glu Pro Asn Phe Val Ala Ala Tyr Asp Ile Gly Leu Phe Ala Tyr
                                         235
                    230
Phe Phe Leu Arg Glu Asn Ala Val Glu His Asp Cys Gly Arg Thr Val
                                     250
                 245
Tyr Ser Arg Val Ala Arg Val Cys Lys Asn Asp Val Gly Gly Arg Phe
                                 265
                                                     270
            260
Leu Leu Glu Asp Thr Trp Thr Thr Phe Met Lys Ala Arg Leu Asn Cys
                                                 285
                             280
 Ser Arg Pro Gly Glu Val Pro Phe Tyr Tyr Asn Glu Leu Gln Ser Ala
                                             300
                         295
 Phe His Leu Pro Glu Gln Asp Leu Ile Tyr Gly Val Phe Thr Thr Asn
```

2760

2820 2880

2940

3000

3060

3120 3180

3237

310

315

Val Asn Ser Ile Ala Ala Ser Ala Val Cys Ala Phe Asn Leu Ser Ala 330 325 Ile Ser Gln Ala Phe Asn Gly Pro Phe Arg Tyr Gln Glu Asn Pro Arg 340 345 Ala Ala Trp Leu Pro Ile Ala Asn Pro Ile Pro Asn Phe Gln Cys Gly 360 Thr Leu Pro Glu Thr Gly Pro Asn Glu Asn Leu Thr Glu Arg Ser Leu 375 380 Gln Asp Ala Gln Arg Leu Phe Leu Met Ser Glu Ala Val Gln Pro Val 390 395 Thr Pro Glu Pro Cys Val Thr Gln Asp Ser Val Arg Phe Ser His Leu 405 410 Val Val Asp Leu Val Gln Ala Lys Asp Thr Leu Tyr His Val Leu Tyr 425 Ile Gly Thr Glu Ser Gly Thr Ile Leu Lys Ala Leu Ser Thr Ala Ser 440 Arg Ser Leu His Gly Cys Tyr Leu Glu Glu Leu His Val Leu Pro Pro 455 460 Gly Arg Arg Glu Pro Leu Arg Ser Leu Arg Ile Leu His Ser Ala Arg 470 475 Ala Leu Phe Val Gly Leu Arg Asp Gly Val Leu Arg Val Pro Leu Glu 485 490 Arg Cys Ala Ala Tyr Arg Ser Gln Gly Ala Cys Leu Gly Ala Arg Asp 505 Pro Tyr Cys Gly Trp Asp Gly Lys Gln Gln Arg Cys Ser Thr Leu Glu 520 Asp Ser Ser Asn Met Ser Leu Trp Thr Gln Asn Ile Thr Ala Cys Pro 535 540 Val Arg Asn Val Thr Arg Asp Gly Gly Phe Gly Pro Trp Ser Pro Trp 550 555 Gln Pro Cys Glu His Leu Asp Gly Asp Asn Ser Gly Ser Cys Leu Cys 570 Arg Ala Arg Ser Cys Asp Ser Pro Arg Pro Arg Cys Gly Gly Leu Asp 585 580 Cys Leu Gly Pro Ala Ile His Ile Ala Asn Cys Ser Arg Asn Gly Ala 600 Trp Thr Pro Trp Ser Ser Trp Ala Leu Cys Ser Thr Ser Cys Gly Ile 615 620 Gly Phe Gln Val Arg Gln Arg Ser Cys Ser Asn Pro Ala Pro Arg His 630 Gly Gly Arg Ile Cys Val Gly Lys Ser Arg Glu Glu Arg Phe Cys Asn 645 650 Glu Asn Thr Pro Cys Pro Val Pro Ile Phe Trp Ala Ser Trp Gly Ser 665 Trp Ser Lys Cys Ser Ser Asn Cys Gly Gly Met Gln Ser Arg Arg 680 Arg Ala Cys Glu Asn Gly Asn Ser Cys Leu Gly Cys Gly Val Glu Phe 695 700 Lys Thr Cys Asn Pro Glu Gly Cys Pro Glu Val Arg Arg Asn Thr Pro 710 715 Trp Thr Pro Trp Leu Pro Val Asn Val Thr Gln Gly Gly Ala Arg Gln 725 730 Glu Gln Arg Phe Arg Phe Thr Cys Arg Ala Pro Leu Ala Asp Pro His 745 Gly Leu Gln Phe Gly Arg Arg Thr Glu Thr Arg Thr Cys Pro Ala 760

```
Asp Gly Ser Gly Ser Cys Asp Thr Asp Ala Leu Val Glu Asp Leu Leu
                      775
                                        780
Arg Ser Gly Ser Thr Ser Pro His Thr Val Ser Gly Gly Trp Ala Ala
                                    795
                  790
Trp Gly Pro Trp Ser Ser Cys Ser Arg Asp Cys Glu Leu Gly Phe Arg
              805
                                810
Val Arg Lys Arg Thr Cys Thr Asn Pro Glu Pro Arg Asn Gly Gly Leu
               825
Pro Cys Val Gly Asp Ala Ala Glu Tyr Gln Asp Cys Asn Pro Gln Ala
                          840
                                           845
Cys Pro Val Arg Gly Ala Trp Ser Cys Trp Thr Ser Trp Ser Pro Cys
                                         860
                      855
Ser Ala Ser Cys Gly Gly Gly His Tyr Gln Arg Thr Arg Ser Cys Thr
                                     875
                  870
Ser Pro Ala Pro Ser Pro Gly Glu Asp Ile Cys Leu Gly Leu His Thr
                                 890
              885
Glu Glu Ala Leu Cys Ala Thr Gln Ala Cys Pro Glu Gly Trp Ser Pro
                             905
           900
Trp Ser Glu Trp Ser Lys Cys Thr Asp Asp Gly Ala Gln Ser Arg Ser
       915 920
Arg His Cys Glu Glu Leu Leu Pro Gly Ser Ser Ala Cys Ala Gly Asn
                                        940
                      935
Ser Ser Gln Ser Arg Pro Cys Pro Tyr Ser Glu Ile Pro Gly Phe Asn
                                     955
                  950
Leu Ile His Leu Val Ala Thr Gly Ile Ser Cys Phe Leu Gly Ser Gly
                                 970
               965
Leu Leu Thr Leu Ala Val Tyr Leu Ser Cys Gln His Cys Gln Arg Gln
                                                990
                              985
           980
Ser Gln Glu Ser Thr Leu Val His Pro Ala Thr Pro Asn His Leu His
                                             1005
                         1000
Tyr Lys Gly Gly Gly Thr Pro Lys Asn Glu Lys Tyr Thr Pro Met Glu
   1010 1015 1020
Phe Lys Thr Leu Asn Lys Asn Asn Leu Ile Pro Asp Asp Arg Ala Asn
1025 - 1030
                        1035 1040
Phe Tyr Pro Leu Gln Gln Thr Asn Val Tyr Thr Thr Thr Tyr Tyr Pro
              1045 1050 1055
Ser Pro Leu Asn Lys His Ser Phe Arg Pro Glu Ala Ser Pro Gly Gln
                             1065
           1060
Arg Cys Phe Pro Asn Ser
        1075
<210> 9
<211> 3456
<212> DNA
<213> homo sapiens
<400> 9
atgccctgtg gcttcagtcc gtctcctgtt gcccaccacc tcgtccctgg gccgcctgat
                                                                  60
accccagccc aacagctaag gtgtggatgg acagtagggg gctggcttct ctcactggtc
                                                                  120
aggggtcttc tcccctgtct gcctcccgga gctaggactg cagaggggcc tatcatggtg
                                                                  180
                                                                  240
cttgcaggcc ccctggctgt ctcgctgttg ctgcccagcc tcacactgct ggtgtcccac
ctctccagct cccaggatgt ctccagtgag cccagcagtg agcagcagct gtgcgccctt
                                                                  300
                                                                  360
agcaagcacc ccaccgtggc ctttgaagac ctgcagccgt gggtctctaa cttcacctac
cctggagccc gggatttctc ccagctggct ttggacccct ccgggracca gctcatcgtg
                                                                  420
 ggagccagga actacctctt cagactcagc cttgccaatg tctctcttct tcaggccaca
                                                                  480
 gagtgggcct ccagtgagga cacgcgccgc tcctgccaaa gcaaagggaa gactgaggag
                                                                  540
```

```
600
gagtgtcaga actacgtgcg agtcctgatc gtcgccggcc ggaaggtgtt catgtgtgga
                                                                       660
accaatgeet ttteececat gtgeaccage agacaggtgg ggaaceteag ceggaetatt
                                                                       720
gagaagatca atggtgtggc ccgctgcccc tatgacccac gccacaactc cacagctgtc
                                                                       780
atctcctccc agggggagct ctatgcagcc acggtcatcg acttctcagg tcgggaccct
gccatctacc gcagectggg cagtgggcca eegettegca etgeccaata taactecaag
                                                                       840
                                                                       900
tggcttaatg agccaaactt cgtggcagcc tatgatattg ggctgtttgc atacttcttc
                                                                       960
ctgcgggaga acgcagtgga gcacgactgt ggacgcaccg tgtactctcg cgtggcccgc
gtgtgcaaga atgacgtggg gggccgattc ctgctggagg acacatggac cacattcatg
                                                                      1020
aaggeeegge teaaetgete eegeeeggge gaggteeeet tetaetataa egagetgeag
                                                                      1080
agtgeettee aettgeerga geaggaeete atetatggag tttteacaae caaegtaaae
                                                                      1140
                                                                      1200
agcatygcgg cttctgctgt ctgcgccttc aacctcagtg ctatctccca ggctttcaat
ggcccatttc gctaccagga gaaccccagg gctgcctggc tccccatagc caaccccatc
                                                                      1260
                                                                      1320
cccaatttcc agtgtggcac cctgcctgag accggtccca acgagaacct gacggagcgc
agcctgcagg acgcgcagcg cctcttcctg atgagcgagg ccgtgcagcc ggtgacaccc
                                                                      1380
gagecetgtg teacceagga cagegtgege tteteacace tegtggtgga eetggtgeag
                                                                      1440
gctaaagaca cgctctacca tgtactctac attggcaccg agtcgggcac catcctgaag
                                                                      1500
                                                                      1560
gcgctgtcca cggcgagccg cagcctccac ggctgctacc tggaggagct gcacgtgctg
                                                                      1620
cccccgggc gccgcgagcc cctgcgcagc ctgcgcatcc tgcacagcgc ccgcgcgctc
                                                                      1680
ttcgtggggc tgagagacgg cgtcctgcgg gtcccactgg agaggtgcgc cgcctaccgc
                                                                      1740
agccaggggg catgcctggg ggcccgggac ccgtactgtg gctgggacgg gaagcagcaa
cgttgcagca cactcgagga cagctccaac atgagcctct ggacccagaa catcaccgcc
                                                                      1800
tgtcctgtgc ggaatgtgac acgggatggg ggcttcggcc catggtcacc atggcaacca
                                                                      1860
tgtgagcact tggatgggga caactcaggc tcttgcctgt gtcgagctcg atcctgtgat
                                                                      1920
                                                                      1980
tecectegae eccgetgtgg gggeettgae tgeetgggge eagecateea eategeeaae
                                                                      2040
tgctccagga atggggcgtg gaccccgtgg tcatcgtggg cgctgtgcag cacgtcctgt
                                                                      2100
ggcatcggct tccaggtccg ccagcgaagt tgcagcaacc ctgctccccg ccacgggggc
                                                                      2160
cgcatctgcg tgggcaagag ccgggaggaa cggttctgta atgagaacac gccttgcccg
                                                                      2220
gtgcccatct tctgggcttc ctggggctcc tggagcaagt gcagcagcaa ctgtggaggg
                                                                      2280
ggcatgcagt cgcggcgtcg ggcctgcgag aacggcaact cctgcctggg ctgcggcgtg
gagttcaaga cgtgcaaccc cgagggctgc cccgaagtgc ggcgcaacac cccctggacg
                                                                      2340
                                                                      2400
ccgtggctgc ccgtgaacgt gacgcagggc ggggcacggc aggagcagcg gttccgcttc
                                                                      2460
acctgccgcg cgccccttgc agacccgcac ggcctgcagt tcggcaggag aaggaccgag
                                                                      2520
acgaggacct gtcccgcgga cggctccggc tcctgcgaca ccgacgccct ggtggaggac
                                                                       2580
ctcctgcgca gcgggagcac ctccccgcac acggtgagcg ggggctgggc cgcctggggc
                                                                       2640
cegtggtegt cetgeteecg ggactgegag etgggettee gegteegeaa gagaacgtge
                                                                       2700
actaacccgg agccccgcaa cgggggcctg ccctgcgtgg gcgatgctgc cgagtaccag
gactgcaacc cccaggcttg cccagttcgg ggtgcttggt cctgctggac ctcatggtct
                                                                       2760
ccatgctcag cttcctgtgg tgggggtcac tatcaacgca cccgttcctg caccagcccc
                                                                       2820
                                                                       2880
 gcaccetece caggtgagga catetgtete gggetgeaca eggaggagge actatgtgee
 acacaggeet geccagaagg etggtegeee tggtetgagt ggagtaagtg eactgaegae
                                                                       2940
ggagcccaga gccgaagccg gcactgtgag gagctcctcc cagggtccag cgcctgtgct
                                                                       3000
 ggaaacagca gccagagccg cccctgcccc tacagcgaga ttcccgtcat cctgccagcc
                                                                       3060
 tecageatgg aggaggeeac eggetgtgea gggtteaate teatecaett ggtggeeacg
                                                                       3120
 ggcatctcct gcttcttggg ctctgggctc ctgaccctag cagtgtacct gtcttgccag
                                                                       3180
 cactgccage gtcagtccca ggagtccaca etggtccate etgccacece caaccatttg
                                                                       3240
                                                                       3300
 cactacaagg gcggaggcac cccgaagaat gaaaagtaca cacccatgga attcaagacc
 ctgaacaaga ataacttgat ccctgatgac agagccaact tctacccatt gcagcagacc
                                                                       3360
 aatgtgtaca cgactactta ctacccaagc cccctgaaca aacacagctt ccggcccgag
                                                                       3420
                                                                       3456
 gcctcacctg gacaacggtg cttccccaac agctga
```

<sup>&</sup>lt;210> 10 <211> 1151

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> homo sapiens

<sup>&</sup>lt;400> 10

```
Met Pro Cys Gly Phe Ser Pro Ser Pro Val Ala His His Leu Val Pro
                                   10
Gly Pro Pro Asp Thr Pro Ala Gln Gln Leu Arg Cys Gly Trp Thr Val
                                25
Gly Gly Trp Leu Leu Ser Leu Val Arg Gly Leu Leu Pro Cys Leu Pro
                           40
Pro Gly Ala Arg Thr Ala Glu Gly Pro Ile Met Val Leu Ala Gly Pro
                       55
Leu Ala Val Ser Leu Leu Leu Pro Ser Leu Thr Leu Leu Val Ser His
                    70
                                       75
Leu Ser Ser Ser Gln Asp Val Ser Ser Glu Pro Ser Ser Glu Gln Gln
                                    90
Leu Cys Ala Leu Ser Lys His Pro Thr Val Ala Phe Glu Asp Leu Gln
                               105
           100
Pro Trp Val Ser Asn Phe Thr Tyr Pro Gly Ala Arg Asp Phe Ser Gln
                                                125
                           120
Leu Ala Leu Asp Pro Ser Gly Asn Gln Leu Ile Val Gly Ala Arg Asn
                        135
Tyr Leu Phe Arg Leu Ser Leu Ala Asn Val Ser Leu Leu Gln Ala Thr
                                        155
                    150
Glu Trp Ala Ser Ser Glu Asp Thr Arg Arg Ser Cys Gln Ser Lys Gly
                                   170
                165
Lys Thr Glu Glu Glu Cys Gln Asn Tyr Val Arg Val Leu Ile Val Ala
                                185
            180
Gly Arg Lys Val Phe Met Cys Gly Thr Asn Ala Phe Ser Pro Met Cys
                            200
Thr Ser Arg Gln Val Gly Asn Leu Ser Arg Thr Ile Glu Lys Ile Asn
                        215
Gly Val Ala Arg Cys Pro Tyr Asp Pro Arg His Asn Ser Thr Ala Val
                                        235
                   230
Ile Ser Ser Gln Gly Glu Leu Tyr Ala Ala Thr Val Ile Asp Phe Ser
                                   250
                245
Gly Arg Asp Pro Ala Ile Tyr Arg Ser Leu Gly Ser Gly Pro Pro Leu
            260
                               265
Arg Thr Ala Gln Tyr Asn Ser Lys Trp Leu Asn Glu Pro Asn Phe Val
                            280
Ala Ala Tyr Asp Ile Gly Leu Phe Ala Tyr Phe Phe Leu Arg Glu Asn
                                           300
                        295
Ala Val Glu His Asp Cys Gly Arg Thr Val Tyr Ser Arg Val Ala Arg
                                        315
                    310
Val Cys Lys Asn Asp Val Gly Gly Arg Phe Leu Leu Glu Asp Thr Trp
                                    330
                325
Thr Thr Phe Met Lys Ala Arg Leu Asn Cys Ser Arg Pro Gly Glu Val
                                345
Pro Phe Tyr Tyr Asn Glu Leu Gln Ser Ala Phe His Leu Pro Glu Gln
                                                365
                            360
        355
 Asp Leu Ile Tyr Gly Val Phe Thr Thr Asn Val Asn Ser Ile Ala Ala
                                            380
                        375
 Ser Ala Val Cys Ala Phe Asn Leu Ser Ala Ile Ser Gln Ala Phe Asn
                                         395
                    390
 Gly Pro Phe Arg Tyr Gln Glu Asn Pro Arg Ala Ala Trp Leu Pro Ile
                                    410
                405
 Ala Asn Pro Ile Pro Asn Phe Gln Cys Gly Thr Leu Pro Glu Thr Gly
                                425
             420
 Pro Asn Glu Asn Leu Thr Glu Arg Ser Leu Gln Asp Ala Gln Arg Leu
                             440
```

```
Phe Leu Met Ser Glu Ala Val Gln Pro Val Thr Pro Glu Pro Cys Val
                       455
Thr Gln Asp Ser Val Arg Phe Ser His Leu Val Val Asp Leu Val Gln
                                      475
                  470
Ala Lys Asp Thr Leu Tyr His Val Leu Tyr Ile Gly Thr Glu Ser Gly
               485
                                  490
Thr Ile Leu Lys Ala Leu Ser Thr Ala Ser Arg Ser Leu His Gly Cys
                              505
Tyr Leu Glu Glu Leu His Val Leu Pro Pro Gly Arg Arg Glu Pro Leu
                                               525
                           520
Arg Ser Leu Arg Ile Leu His Ser Ala Arg Ala Leu Phe Val Gly Leu
                                           540
                       535
Arg Asp Gly Val Leu Arg Val Pro Leu Glu Arg Cys Ala Ala Tyr Arg
                                       555
                   550
Ser Gln Gly Ala Cys Leu Gly Ala Arg Asp Pro Tyr Cys Gly Trp Asp
                                  570
               565
Gly Lys Gln Gln Arg Cys Ser Thr Leu Glu Asp Ser Ser Asn Met Ser
                               585
Leu Trp Thr Gln Asn Ile Thr Ala Cys Pro Val Arg Asn Val Thr Arg
                           600
       595
Asp Gly Gly Phe Gly Pro Trp Ser Pro Trp Gln Pro Cys Glu His Leu
                                           620
                       615
Asp Gly Asp Asn Ser Gly Ser Cys Leu Cys Arg Ala Arg Ser Cys Asp
                                       635
                    630
Ser Pro Arg Pro Arg Cys Gly Gly Leu Asp Cys Leu Gly Pro Ala Ile
                                    650
                645
His Ile Ala Asn Cys Ser Arg Asn Gly Ala Trp Thr Pro Trp Ser Ser
                               665
Trp Ala Leu Cys Ser Thr Ser Cys Gly Ile Gly Phe Gln Val Arg Gln
                           680
Arg Ser Cys Ser Asn Pro Ala Pro Arg His Gly Gly Arg Ile Cys Val
                                           700
                       695
Gly Lys Ser Arg Glu Glu Arg Phe Cys Asn Glu Asn Thr Pro Cys Pro
                            715
                    710
Val Pro Ile Phe Trp Ala Ser Trp Gly Ser Trp Ser Lys Cys Ser Ser
                                   730
               725
Asn Cys Gly Gly Met Gln Ser Arg Arg Arg Ala Cys Glu Asn Gly
                                745
            740
Asn Ser Cys Leu Gly Cys Gly Val Glu Phe Lys Thr Cys Asn Pro Glu
                            760
Gly Cys Pro Glu Val Arg Arg Asn Thr Pro Trp Thr Pro Trp Leu Pro
                                            780
                        775
Val Asn Val Thr Gln Gly Gly Ala Arg Gln Glu Gln Arg Phe Arg Phe
                                        795
Thr Cys Arg Ala Pro Leu Ala Asp Pro His Gly Leu Gln Phe Gly Arg
                                   810
                805
Arg Arg Thr Glu Thr Arg Thr Cys Pro Ala Asp Gly Ser Gly Ser Cys
                                                    830
                                825
            820
 Asp Thr Asp Ala Leu Val Glu Asp Leu Leu Arg Ser Gly Ser Thr Ser
                            840
 Pro His Thr Val Ser Gly Gly Trp Ala Ala Trp Gly Pro Trp Ser Ser
                                            860
                        855
 Cys Ser Arg Asp Cys Glu Leu Gly Phe Arg Val Arg Lys Arg Thr Cys
                                        875
                    870
 Thr Asn Pro Glu Pro Arg Asn Gly Gly Leu Pro Cys Val Gly Asp Ala
                                    890
                 885
```

```
Ala Glu Tyr Gln Asp Cys Asn Pro Gln Ala Cys Pro Val Arg Gly Ala
                              905
           900
Trp Ser Cys Trp Thr Ser Trp Ser Pro Cys Ser Ala Ser Cys Gly Gly
                                            925
                       920
       915
Gly His Tyr Gln Arg Thr Arg Ser Cys Thr Ser Pro Ala Pro Ser Pro
                   935
                                     940
Gly Glu Asp Ile Cys Leu Gly Leu His Thr Glu Glu Ala Leu Cys Ala
                       955
                  950
945
Thr Gln Ala Cys Pro Glu Gly Trp Ser Pro Trp Ser Glu Trp Ser Lys
                                  970
               965
Cys Thr Asp Asp Gly Ala Gln Ser Arg Ser Arg His Cys Glu Glu Leu
                              985
           980
Leu Pro Gly Ser Ser Ala Cys Ala Gly Asn Ser Ser Gln Ser Arg Pro
                          1000
Cys Pro Tyr Ser Glu Ile Pro Val Ile Leu Pro Ala Ser Ser Met Glu
                     1015
                                        1020
    1010
Glu Ala Thr Gly Cys Ala Gly Phe Asn Leu Ile His Leu Val Ala Thr
                  1030
                                    1035
Gly Ile Ser Cys Phe Leu Gly Ser Gly Leu Leu Thr Leu Ala Val Tyr
               1045 1050
Leu Ser Cys Gln His Cys Gln Arg Gln Ser Gln Glu Ser Thr Leu Val
                                                1070
                              1065
           1060
His Pro Ala Thr Pro Asn His Leu His Tyr Lys Gly Gly Gly Thr Pro
                                             1085
                          1080
Lys Asn Glu Lys Tyr Thr Pro Met Glu Phe Lys Thr Leu Asn Lys Asn
                                         1100
                       1095
Asn Leu Ile Pro Asp Asp Arg Ala Asn Phe Tyr Pro Leu Gln Gln Thr
                   1110
                                      1115
Asn Val Tyr Thr Thr Tyr Tyr Pro Ser Pro Leu Asn Lys His Ser
                                 1130
               1125
Phe Arg Pro Glu Ala Ser Pro Gly Gln Arg Cys Phe Pro Asn Ser
                              1145
            1140
<210> 11
<211> 3411
<212> DNA
<213> homo sapiens
<400> 11
 atgccctgtg gcttcagtcc gtctcctgtt gcccaccacc tcgtccctgg gccgcctgat
                                                                   60
                                                                   120
 accccagccc aacagctaag gtgtggatgg acagtagggg gctggcttct ctcactggtc
 aggggtette teccetgtet geeteeegga getaggaetg cagaggggee tateatggtg
                                                                   180
 cttgcaggcc ccctggctgt ctcgctgttg ctgcccagcc tcacactgct ggtgtcccac
                                                                   240
 ctctccagct cccaggatgt ctccagtgag cccagcagtg agcagcagct gtgcgccctt
                                                                   300
 agcaagcacc ccaccgtggc ctttgaagac ctgcagccgt gggtctctaa cttcacctac
                                                                   360
 cctggagccc gggatttctc ccagctggct ttggacccct ccgggracca gctcatcgtg
                                                                   420
                                                                   480
 ggagccagga actacctctt cagactcagc cttgccaatg tctctcttct tcaggccaca
 gagtgggcct ccagtgagga cacgcgccgc tcctgccaaa gcaaagggaa gactgaggag
                                                                   540
                                                                   600
 gagtgtcaga actacgtgcg agtcctgatc gtcgccggcc ggaaggtgtt catgtgtgga
 accaatgeet ttteececat gtgeaccage agacaggtgg ggaaceteag ceggaetatt
                                                                   660
 gagaagatca atggtgtggc ccgctgcccc tatgacccac gccacaactc cacagctgtc
                                                                   720
 atctcctccc agggggagct ctatgcagcc acggtcatcg acttctcagg tcgggaccct
                                                                   780
 gccatctacc gcagcctggg cagtgggcca ccgcttcgca ctgcccaata taactccaag
                                                                   840
 tggcttaatg agccaaactt cgtggcagcc tatgatattg ggctgtttgc atacttcttc
                                                                   900
 ctgcgggaga acgcagtgga gcacgactgt ggacgcaccg tgtactctcg cgtggcccgc
                                                                   960
 gtgtgcaaga atgacgtggg gggccgattc ctgctggagg acacatggac cacattcatg
                                                                  1020
```

```
aaggcccggc tcaactgctc ccgcccgggc gaggtcccct tctactataa cgagctgcag
                                                                     1080
agtgccttcc acttgccrga gcaggacctc atctatggag ttttcacaac caacgtaaac
                                                                      1140
agcatygcgg cttctgctgt ctgcgccttc aacctcagtg ctatctccca ggctttcaat
                                                                      1200
ggcccatttc gctaccagga gaaccccagg gctgcctggc tccccatagc caaccccate
                                                                      1260
                                                                      1320
cccaatttcc agtgtggcac cctgcctgag accggtccca acgagaacct gacggagcgc
agcctgcagg acgcgcagcg cctcttcctg atgagcgagg ccgtgcagcc ggtgacaccc
                                                                      1380
                                                                      1440
gagecetgtg teacceagga cagegtgege tteteacace tegtggtgga cetggtgcag
gctaaagaca cgctctacca tgtactctac attggcaccg agtcgggcac catcctgaag
                                                                      1500
gegetgteca eggegageeg eageeteeae ggetgetaee tggaggaget geaegtgetg
                                                                      1560
cccccgggc gccgcgagcc cctgcgcagc ctgcgcatcc tgcacagcgc ccgcgcgctc
                                                                      1620
                                                                      1680
ttcgtggggc tgagagacgg cgtcctgcgg gtcccactgg agaggtgcgc cgcctaccgc
                                                                      1740
agccaggggg catgcctggg ggcccgggac ccgtactgtg gctgggacgg gaagcagcaa
                                                                      1800
cgttgcagca cactcgagga cagctccaac atgagcctct ggacccagaa catcaccgcc
tgtcctgtgc ggaatgtgac acgggatggg ggcttcggcc catggtcacc atggcaacca
                                                                      1860
                                                                      1920
tgtgagcact tggatgggga caactcaggc tcttgcctgt gtcgagctcg atcctgtgat
tecectegae ecceetegteg gegeettegae tecetegege cagecateca categecaae
                                                                      1980
                                                                      2040
tgctccagga atggggcgtg gaccccgtgg tcatcgtggg cgctgtgcag cacgtcctgt
                                                                      2100
ggcatcggct tccaggtccg ccagcgaagt tgcagcaacc ctgctccccg ccacgggggc
                                                                      2160
cgcatctgcg tgggcaagag ccgggaggaa cggttctgta atgagaacac gccttgcccg
                                                                      2220
gtgcccatct tctgggcttc ctggggctcc tggagcaagt gcagcagcaa ctgtggaggg
ggcatgcagt cgcggcgtcg ggcctgcgag aacggcaact cctgcctggg ctgcggcgtg
                                                                      2280
                                                                      2340
gagttcaaga cgtgcaaccc cgagggctgc cccgaagtgc ggcgcaacac cccctggacg
ccgtggctgc ccgtgaacgt gacgcagggc ggggcacggc aggagcagcg gttccgcttc
                                                                      2400
acctgccgcg cgccccttgc agacccgcac ggcctgcagt tcggcaggag aaggaccgag
                                                                      2460
                                                                      2520
acgaggacct gtcccgcgga cggctccggc tcctgcgaca ccgacgccct ggtggaggac
ctcctgcgca gcgggagcac ctccccgcac acggtgagcg ggggctgggc cgcctggggc
                                                                      2580
                                                                      2640
ccgtggtcgt cctgctcccg ggactgcgag ctgggcttcc gcgtccgcaa gagaacgtgc
                                                                      2700
actaaccegg ageceegeaa egggggeetg eeetgegtgg gegatgetge egagtaecag
                                                                      2760
gactgcaacc cccaggettg cccagttcgg ggtgcttggt cctgctggac ctcatggtct
ccatgctcag cttcctgtgg tgggggtcac tatcaacgca cccgttcctg caccagcccc
                                                                      2820
                                                                      2880
gcaccetece caggtgagga catetgtete gggetgeaca eggaggagge actatgtgee
                                                                      2940
acacaggeet geccagaagg etggtegeee tggtetgagt ggagtaagtg eactgaegae
                                                                      3000
ggagcccaga gccgaagccg gcactgtgag gagctcctcc cagggtccag cgcctgtgct
ggaaacagca gccagagccg cccctgcccc tacagcgaga ttcccgggtt caatctcatc
                                                                      3060
                                                                      3120
cacttggtgg ccacgggcat ctcctgcttc ttgggctctg ggctcctgac cctagcagtg
                                                                      3180
tacctgtctt gccagcactg ccagcgtcag tcccaggagt ccacactggt ccatcctgcc
acccccaacc atttgcacta caagggcgga ggcaccccga agaatgaaaa gtacacaccc
                                                                      3240
atggaattca agaccctgaa caagaataac ttgatccctg atgacagagc caacttctac
                                                                      3300
                                                                      3360
ccattgcagc agaccaatgt gtacacgact acttactacc caagccccct gaacaaacac
agcttccggc ccgaggcctc acctggacaa cggtgcttcc ccaacagctg a
                                                                      3411
```

<210> 12 <211> 1136

<212> PRT <213> homo sapiens

<400> 12

 Met
 Pro
 Cys
 Gly
 Phe
 Ser
 Pro
 Ser
 Pro
 Val
 Ala
 His
 Leu
 Val
 Pro

 Gly
 Pro
 Pro
 Asp
 Thr
 Pro
 Ala
 Gln
 Gln
 Leu
 Arg
 Cys
 Gly
 Trp
 Thr
 Val

 Gly
 Gly
 Trp
 Leu
 Leu
 Ser
 Leu
 Val
 Arg
 Gly
 Leu
 Pro
 Cys
 Leu
 Pro

 Pro
 Gly
 Ala
 Arg
 Thr
 Ala
 Gly
 Pro
 Ile
 Met
 Val
 Leu
 Ala
 Gly
 Pro

 Fro
 Gly
 Ala
 Arg
 Thr
 Ala
 Gly
 Pro
 Ile
 Met
 Val
 Leu
 Ala
 Gly
 Pro

 Brown
 Arg
 Arg
 Fro
 Arg
 Pro
 Ile
 Met
 Val
 Leu
 Ala
 Gly
 Pro
 Gly
 Fro
 Ile

65					70					75			_		80
Leu S				85					90					95	
Leu (			100					105					110		
Pro 7		Val 115	Ser	Asn	Phe	Thr	Tyr 120	Pro	Gly	Ala	Arg	Asp 125	Phe	Ser	Gln
Leu A			Asp	Pro	Ser	Gly 135	Asn	Gln	Leu	Ile	Val 140	Gly	Ala	Arg	Asn
Tyr 1	Leu	Phe	Arg	Leu	Ser 150	Leu	Ala	Asn	Val	Ser 155	Leu	Leu	Gln	Ala	Thr 160
Glu '				165					170					175	
Lys '			180					185					190		
Gly .		195					200					205		-	
Thr	210					215					220				
Gly 225					230					235					240
				245					250	Thr				255	
Gly			260					265					270		
		275					280			Asn		285			
	290					295				Phe	300				
305					310					Tyr 315					320
				325					330					335	
			340					345					350		Val
		355					360					365			Gln
	370					375	,				380				Ala
385					390	i				395					Asn 400
				405					410	)				415	
			420	ı				425	5				430	)	Gly
		435	5				440	)				445			Leu
	450					455	5				460	)			val
465					470	)				475	5				480
				485	5				490	)				495	
			500	)				50!	5				510	)	y Cys
Tyr	Leu	ı Glı	ı Glu	ı Lev	ı His	s Vai	l Le	ı Pro	o Pro	o Gly	/ Arc	y Arc	g Glu	ı Pro	Leu

		515					520					525			
Arg	Ser 530	Leu	Arg	Ile	Leu	His 535	Ser	Ala	Arg	Ala	Leu 540	Phe	Val	Gly	Leu
Arg 545	Asp	Gly	Val	Leu	Arg 550	Val	Pro	Leu	Glu	Arg 555	Cys	Ala	Ala	Tyr	Arg 560
Ser	Gln	Gly	Ala	Суз 565	Leu	Gly	Ala	Arg	Asp 570	Pro	Tyr	Cys	Gly	Trp 575	Asp
Gly	Lys	Gln	Gln 580	Arg	Cys	Ser	Thr	Leu 585	Glu	Asp	Ser	Ser	Asn 590	Met	Ser
Leu	Trp	Thr 595	Gln	Asn	Ile	Thr	Ala 600	Cys	Pro	Val	Arg	Asn 605	Val	Thr	Arg
_	610				Pro	615					620				
625					Gly 630					635					640
				645	Cys				650					655	
			660		Ser			665					670		
_		675			Thr		680					685			
	690				Pro	695					700				
705					Glu 710					715					720
				725	Ala				730					735	
			740		Met			745					750		
		755			Cys		760			,		765			
	770				Arg	775					780				
785					Gly 790					795					800
				805					810					815	
			820					825					830		Cys
		835					840					845	•		Ser
	850	)				855					860				Ser
865					870					875	i				880
				885	· •				890	)				895	
			900	)				905	•				910	)	Ala
		915	5				920					925	5		Gly
	930	)				935	,				940	)			r Pro
945	5				950	1				955	5				960
Thr	Gli	n Ala	a Cys	s Pro	o Glu	ı Gly	Trp	Ser	Pro	o Tr	ser ser	GII	ı 'l'Y	, sei	. Lys

```
970
               965
Cys Thr Asp Asp Gly Ala Gln Ser Arg Ser Arg His Cys Glu Glu Leu
          980 985
Leu Pro Gly Ser Ser Ala Cys Ala Gly Asn Ser Ser Gln Ser Arg Pro
       995
                         1000 1005
Cys Pro Tyr Ser Glu Ile Pro Gly Phe Asn Leu Ile His Leu Val Ala
                       1015
                                           1020
Thr Gly Ile Ser Cys Phe Leu Gly Ser Gly Leu Leu Thr Leu Ala Val
                   1030
                                       1035
Tyr Leu Ser Cys Gln His Cys Gln Arg Gln Ser Gln Glu Ser Thr Leu
               1045
                                   1050
                                                       1055
Val His Pro Ala Thr Pro Asn His Leu His Tyr Lys Gly Gly Thr
           1060
                               1065
                                                   1070
Pro Lys Asn Glu Lys Tyr Thr Pro Met Glu Phe Lys Thr Leu Asn Lys
                           1080
                                               1085
Asn Asn Leu Ile Pro Asp Asp Arg Ala Asn Phe Tyr Pro Leu Gln Gln
                       1095
                                           1100
Thr Asn Val Tyr Thr Thr Tyr Tyr Pro Ser Pro Leu Asn Lys His
                   1110
                                       1115
Ser Phe Arg Pro Glu Ala Ser Pro Gly Gln Arg Cys Phe Pro Asn Ser
                                                       1135
               1125
                                   1130
<210> 13
<211> 2865
<212> DNA
<213> homo sapiens
<400> 13
                                                                      60
atgtgtggaa ccaatgcctt ttcccccatg tgcaccagca gacaggtggg gaacctcagc
                                                                     120
cggactattg agaagatcaa tggtgtggcc cgctgcccct atgacccacg ccacaactcc
                                                                     180
acagetgtea teteeteeca gggggagete tatgeageea eggteatega etteteaggt
                                                                     240
egggaeeetg ceatetaeeg eageetggge agtgggeeae egettegeae tgeecaatat
                                                                     300
aactccaagt ggcttaatga gccaaacttc gtggcagcct atgatattgg gctgtttgca
tacttcttcc tgcgggagaa cgcagtggag cacgactgtg gacgcaccgt gtactctcgc
                                                                     360
                                                                     420
gtggcccgcg tgtgcaagaa tgacgtgggg ggccgattcc tgctggagga cacatggacc
acattcatga aggcccggct caactgctcc cgcccgggcg aggtcccctt ctactataac
                                                                     480
                                                                     540
gagetgeaga gtgeetteea ettgeergag eaggaeetea tetatggagt ttteacaace
                                                                     600
aacgtaaaca gcatygcggc ttctgctgtc tgcgccttca acctcagtgc tatctcccag
gctttcaatg gcccatttcg ctaccaggag aaccccaggg ctgcctggct ccccatagcc
                                                                     660
aaccccatcc ccaatttcca gtgtggcacc ctgcctgaga ccggtcccaa cgagaacctg
                                                                     720
acggagegea geetgeagga egegeagege etetteetga tgagegagge egtgeageeg
                                                                     780
                                                                     840
gtgacacccg agccctgtgt cacccaggac agcgtgcgct tctcacacct cgtggtggac
ctggtgcagg ctaaagacac gctctaccat gtactctaca ttggcaccga gtcgggcacc
                                                                     900
atcctgaagg cgctgtccac ggcgagccgc agcctccacg gctgctacct ggaggagctg
                                                                     960
cacqtgctgc cccccgggcg ccgcgagccc ctgcgcagcc tgcgcatcct gcacagcgcc
                                                                    1020
cgcgcgctct tcgtggggct gagagacggc gtcctgcggg tcccactgga gaggtgcgcc
                                                                    1080
gcctaccgca gccagggggc atgcctgggg gcccgggacc cgtactgtgg ctgggacggg
                                                                    1140
aagcagcaac gttgcagcac actcgaggac agctccaaca tgagcctctg gacccagaac
                                                                    1200
                                                                    1260
atcaccgcct gtcctgtgcg gaatgtgaca cgggatgggg gcttcggccc atggtcacca
                                                                    1320
tggcaaccat gtgagcactt ggatggggac aactcaggct cttgcctgtg tcgagctcga
tectgtgatt eccetegace eegetgtggg ggeettgact geetggggee ageeateeae
                                                                    1380
ategecaact getecaggaa tggggegtgg acceegtggt categtggge getgtgeage
                                                                    1440
                                                                    1500
acgtectgtg gcatcggctt ccaggtccgc cagcgaagtt gcagcaaccc tgctccccgc
                                                                    1560
cacgggggcc gcatctgcgt gggcaagagc cgggaggaac ggttctgtaa tgagaacacg
                                                                    1620
ccttgcccgg tgcccatctt ctgggcttcc tggggctcct ggagcaagtg cagcagcaac
```

tgtggagggg gcatgcagtc gcggcgtcgg gcctgcgaga acggcaactc ctgcctgggc

```
tgcggcgtgg agttcaagac gtgcaacccc gagggctgcc ccgaagtgcg gcgcaacacc
                                                                      1740
ccctggacgc cgtggctgcc cgtgaacgtg acgcagggcg gggcacggca ggagcagcgg
                                                                      1800
                                                                      1860
ttccgcttca cctgccgcgc gccccttgca gacccgcacg gcctgcagtt cggcaggaga
aggaccgaga cgaggacctg tcccgcggac ggctccggct cctgcgacac cgacgccctg
                                                                      1920
                                                                      1980
gtggaggacc tcctgcgcag cgggagcacc tccccgcaca cggtgagcgg gggctgggcc
gcctggggcc cgtggtcgtc ctgctcccgg gactgcgagc tgggcttccg cgtccgcaag
                                                                      2040
agaacgtgca ctaacccgga gccccgcaac gggggcctgc cctgcgtggg cgatgctgcc
                                                                      2100
                                                                      2160
gagtaccagg actgcaaccc ccaggcttgc ccagttcggg gtgcttggtc ctgctggacc
tcatggtctc catgctcagc ttcctgtggt gggggtcact atcaacgcac ccgttcctgc
                                                                      2220
                                                                      2280
accageceeg caeceteece aggtgaggae atetgteteg ggetgeacae ggaggaggea
                                                                      2340
ctatgtgcca cacaggcctg cccagaaggc tggtcgccct ggtctgagtg gagtaagtgc
                                                                      2400
actgacgacg gagcccagag ccgaagccgg cactgtgagg agctcctccc agggtccagc
geetgtgetg gaaacageag eeagageege eeetgeeeet acagegagat teeegteate
                                                                      2460
ctgccagcct ccagcatgga ggaggccacc ggctgtgcag ggttcaatct catccacttg
                                                                      2520
gtggccacgg gcatctcctg cttcttgggc tctgggctcc tgaccctagc agtgtacctg
                                                                      2580
                                                                      2640
tettgecage actgecageg teagteceag gagtecaeae tggtecatee tgecaeeeee
aaccatttgc actacaaggg cggaggcacc ccgaagaatg aaaagtacac acccatggaa
                                                                      2700
ttcaagaccc tgaacaagaa taacttgatc cctgatgaca gagccaactt ctacccattg
                                                                      2760
cagcagacca atgtgtacac gactacttac tacccaagcc ccctgaacaa acacagcttc
                                                                      2820
                                                                      2865
cggcccgagg cctcacctgg acaacggtgc ttccccaaca gctga
```

<210> 14 <211> 954 <212> PRT

<213> homo sapiens

<400> 14

Met Cys Gly Thr Asn Ala Phe Ser Pro Met Cys Thr Ser Arg Gln Val 15 5 Gly Asn Leu Ser Arg Thr Ile Glu Lys Ile Asn Gly Val Ala Arg Cys 25 Pro Tyr Asp Pro Arg His Asn Ser Thr Ala Val Ile Ser Ser Gln Gly 40 Glu Leu Tyr Ala Ala Thr Val Ile Asp Phe Ser Gly Arg Asp Pro Ala 60 55 Ile Tyr Arg Ser Leu Gly Ser Gly Pro Pro Leu Arg Thr Ala Gln Tyr 75 70 Asn Ser Lys Trp Leu Asn Glu Pro Asn Phe Val Ala Ala Tyr Asp Ile 85 90 Gly Leu Phe Ala Tyr Phe Phe Leu Arg Glu Asn Ala Val Glu His Asp 105 100 Cys Gly Arg Thr Val Tyr Ser Arg Val Ala Arg Val Cys Lys Asn Asp 125 120 115 Val Gly Gly Arg Phe Leu Leu Glu Asp Thr Trp Thr Thr Phe Met Lys 140 135 Ala Arg Leu Asn Cys Ser Arg Pro Gly Glu Val Pro Phe Tyr Tyr Asn 150 155 145 Glu Leu Gln Ser Ala Phe His Leu Pro Glu Gln Asp Leu Ile Tyr Gly 165 170 Val Phe Thr Thr Asn Val Asn Ser Ile Ala Ala Ser Ala Val Cys Ala 190 185 180 Phe Asn Leu Ser Ala Ile Ser Gln Ala Phe Asn Gly Pro Phe Arg Tyr 200 Gln Glu Asn Pro Arg Ala Ala Trp Leu Pro Ile Ala Asn Pro Ile Pro 220 215 Asn Phe Gln Cys Gly Thr Leu Pro Glu Thr Gly Pro Asn Glu Asn Leu

225					230					235					240
	Glu	Arg	Ser	Leu 245		Asp	Ala	Gln	Arg 250	Leu	Phe	Leu	Met	Ser 255	Glu
			Pro 260					265					270		
		275	His				280					285			
	290		Leu			295					300				
305			Ala		310					315					320
			Pro	325					330					335	
			Ala 340					345					350		
		355	Leu				360					365			
	370		Arg			375					380				
385			Leu		390					395					400
			Cys	405					410					415	
			Pro 420					425					430		
		435	Leu				440					445			
	450		Leu			455					460				
465			Gly		470					475					480
			Gly	485					490					495	
			Arg 500					505					510		
		515	Cys				520					525			
	530					535					540				Gly
545					550					555					Gly 560
			Glu	565					570					575	
			580					585					590		Gln
		595					600					605			Pro
	610					615					620				Thr
625					630					635					Leu 640
				645	<b>,</b>				650					655	Ser
			660					665					670		Cys
Glu	Lev	Gly	Phe	Arg	y Val	Arg	Lys	Arg	Thr	Cys	Thr	Asn	Pro	Glu	Pro

```
675
                            680
Arg Asn Gly Gly Leu Pro Cys Val Gly Asp Ala Ala Glu Tyr Gln Asp
                                            700
                        695
Cys Asn Pro Gln Ala Cys Pro Val Arg Gly Ala Trp Ser Cys Trp Thr
                                        715
                    710
Ser Trp Ser Pro Cys Ser Ala Ser Cys Gly Gly Gly His Tyr Gln Arg
                                    730
                725
Thr Arg Ser Cys Thr Ser Pro Ala Pro Ser Pro Gly Glu Asp Ile Cys
                                745
                                                    750
            740
Leu Gly Leu His Thr Glu Glu Ala Leu Cys Ala Thr Gln Ala Cys Pro
                            760
Glu Gly Trp Ser Pro Trp Ser Glu Trp Ser Lys Cys Thr Asp Asp Gly
                                            780
                        775
Ala Gln Ser Arg Ser Arg His Cys Glu Glu Leu Leu Pro Gly Ser Ser
                                        795
                    790
Ala Cys Ala Gly Asn Ser Ser Gln Ser Arg Pro Cys Pro Tyr Ser Glu
                                    810
                805
Ile Pro Val Ile Leu Pro Ala Ser Ser Met Glu Glu Ala Thr Gly Cys
                                                     830
                                825
            820
Ala Gly Phe Asn Leu Ile His Leu Val Ala Thr Gly Ile Ser Cys Phe
                            840
Leu Gly Ser Gly Leu Leu Thr Leu Ala Val Tyr Leu Ser Cys Gln His
                                             860
                        855
    850
Cys Gln Arg Gln Ser Gln Glu Ser Thr Leu Val His Pro Ala Thr Pro
                                         875
                    870
Asn His Leu His Tyr Lys Gly Gly Gly Thr Pro Lys Asn Glu Lys Tyr
                                                         895
                                     890
                885
Thr Pro Met Glu Phe Lys Thr Leu Asn Lys Asn Asn Leu Ile Pro Asp
                                                     910
                                905
Asp Arg Ala Asn Phe Tyr Pro Leu Gln Gln Thr Asn Val Tyr Thr Thr
                                                 925
                            920
Thr Tyr Tyr Pro Ser Pro Leu Asn Lys His Ser Phe Arg Pro Glu Ala
                                             940
                       935
Ser Pro Gly Gln Arg Cys Phe Pro Asn Ser
                     950
<210> 15
<211> 2820
<212> DNA
<213> homo sapiens
<400> 15
                                                                         60
atgtgtggaa ccaatgcctt ttcccccatg tgcaccagca gacaggtggg gaacctcagc
cggactattg agaagatcaa tggtgtggcc cgctgcccct atgacccacg ccacaactcc
                                                                        120
acagctgtca tctcctccca gggggagctc tatgcagcca cggtcatcga cttctcaggt
                                                                        180
                                                                        240
cgggaccctg ccatctaccg cagcctgggc agtgggccac cgcttcgcac tgcccaatat
aactccaagt ggcttaatga gccaaacttc gtggcagcct atgatattgg gctgtttgca
                                                                        300
                                                                        360
tacttcttcc tgcgggagaa cgcagtggag cacgactgtg gacgcaccgt gtactctcgc
 gtggcccgcg tgtgcaagaa tgacgtgggg ggccgattcc tgctggagga cacatggacc
                                                                        420
 acattcatga aggcccggct caactgctcc cgcccgggcg aggtcccctt ctactataac
                                                                        480
 gagetgeaga gtgeetteea ettgeergag eaggaeetea tetatggagt ttteaeaace
                                                                        540
 aacgtaaaca gcatygcggc ttctgctgtc tgcgccttca acctcagtgc tatctcccag
                                                                        600
 gctttcaatg gcccatttcg ctaccaggag aaccccaggg ctgcctggct ccccatagcc
                                                                        660
 aaccccatcc ccaatttcca gtgtggcacc ctgcctgaga ccggtcccaa cgagaacctg
                                                                        720
                                                                        780
 acggagcgca gcctgcagga cgcgcagcgc ctcttcctga tgagcgaggc cgtgcagccg
 gtgacacccg agccctgtgt cacccaggac agcgtgcgct tctcacacct cgtggtggac
                                                                        840
```

```
900
ctggtgcagg ctaaagacac gctctaccat gtactctaca ttggcaccga gtcgggcacc
                                                                      960
affectgaagg egetgteeae ggegageege ageeteeaeg getgetaeet ggaggagetg
                                                                     1020
cacgtgctgc ccccgggcg ccgcgagccc ctgcgcagcc tgcgcatcct gcacagcgcc
cgcgcgctct tcgtggggct gagagacggc gtcctgcggg tcccactgga gaggtgcgcc
                                                                     1080
                                                                     1140
gcctaccgca gccagggggc atgcctgggg gcccgggacc cgtactgtgg ctgggacggg
                                                                     1200
aagcagcaac gttgcagcac actcgaggac agctccaaca tgagcctctg gacccagaac
                                                                     1260
atcaccgcct gtcctgtgcg gaatgtgaca cgggatgggg gcttcggccc atggtcacca
tggcaaccat gtgagcactt ggatggggac aactcaggct cttgcctgtg tcgagctcga
                                                                     1320
                                                                     1380
tcctgtgatt cccctcgacc ccgctgtggg ggccttgact gcctggggcc agccatccac
                                                                     1440
atcgccaact gctccaggaa tggggcgtgg accccgtggt catcgtgggc gctgtgcagc
                                                                     1500
acgtcctgtg gcatcggctt ccaggtccgc cagcgaagtt gcagcaaccc tgctccccgc
                                                                     1560
cacgggggcc gcatctgcgt gggcaagagc cgggaggaac ggttctgtaa tgagaacacg
                                                                     1620
cettgecegg tgeccatett etgggettee tggggeteet ggagcaagtg cagcagcaac
tgtggagggg gcatgcagtc gcggcgtcgg gcctgcgaga acggcaactc ctgcctgggc
                                                                     1680
tgcggcgtgg agttcaagac gtgcaacccc gagggctgcc ccgaagtgcg gcgcaacacc
                                                                     1740
ccctggacgc cgtggctgcc cgtgaacgtg acgcagggcg gggcacggca ggagcagcgg
                                                                     1800
                                                                     1860
ttccgcttca cctgccgcgc gccccttgca gacccgcacg gcctgcagtt cggcaggaga
aggaccgaga cgaggacctg tcccgcggac ggctccggct cctgcgacac cgacgccctg
                                                                     1920
                                                                     1980
gtggaggacc tcctgcgcag cgggagcacc tccccgcaca cggtgagcgg gggctgggcc
gcctggggcc cgtggtcgtc ctgctcccgg gactgcgagc tgggcttccg cgtccgcaag
                                                                     2040
agaacgtgca ctaacccgga gccccgcaac gggggcctgc cctgcgtggg cgatgctgcc
                                                                     2100
gagtaccagg actgcaaccc ccaggcttgc ccagttcggg gtgcttggtc ctgctggacc
                                                                     2160
tcatggtctc catgctcagc ttcctgtggt gggggtcact atcaacgcac ccgttcctgc
                                                                     2220
                                                                     2280
accageceeg caeeeteece aggtgaggae atetgteteg ggetgeacae ggaggaggea
ctatgtgcca cacaggcctg cccagaaggc tggtcgccct ggtctgagtg gagtaagtgc
                                                                     2340
                                                                     2400
actgacgacg gagcccagag ccgaagccgg cactgtgagg agctcctccc agggtccagc
gcctgtgctg gaaacagcag ccagagccgc ccctgcccct acagcgagat tcccgggttc
                                                                     2460
aatctcatcc acttggtggc cacgggcatc tcctgcttct tgggctctgg gctcctgacc
                                                                     2520
ctagcagtgt acctgtcttg ccagcactgc cagcgtcagt cccaggagtc cacactggtc
                                                                     2580
catcctgcca ccccaacca tttgcactac aagggcggag gcaccccgaa gaatgaaaag
                                                                     2640
                                                                     2700
tacacaccca tggaattcaa gaccctgaac aagaataact tgatccctga tgacagagcc
aacttctacc cattgcagca gaccaatgtg tacacgacta cttactaccc aagccccctg
                                                                     2760
aacaaacaca getteeggee egaggeetea eetggacaae ggtgetteee caacagetga
                                                                     2820
<210> 16
<211> 939
```

<212> PRT

<213> homo sapiens

<400> 16

Met Cys Gly Thr Asn Ala Phe Ser Pro Met Cys Thr Ser Arg Gln Val 1 5 10 15 Gly Asn Leu Ser Arg Thr Ile Glu Lys Ile Asn Gly Val Ala Arg Cys 25 20 Pro Tyr Asp Pro Arg His Asn Ser Thr Ala Val Ile Ser Ser Gln Gly 40 45 Glu Leu Tyr Ala Ala Thr Val Ile Asp Phe Ser Gly Arg Asp Pro Ala 55 60 Ile Tyr Arg Ser Leu Gly Ser Gly Pro Pro Leu Arg Thr Ala Gln Tyr 70 75 Asn Ser Lys Trp Leu Asn Glu Pro Asn Phe Val Ala Ala Tyr Asp Ile 90 85 Gly Leu Phe Ala Tyr Phe Phe Leu Arg Glu Asn Ala Val Glu His Asp 105 Cys Gly Arg Thr Val Tyr Ser Arg Val Ala Arg Val Cys Lys Asn Asp 115 120

```
Val Gly Gly Arg Phe Leu Leu Glu Asp Thr Trp Thr Thr Phe Met Lys
                        135
                                            140
Ala Arg Leu Asn Cys Ser Arg Pro Gly Glu Val Pro Phe Tyr Tyr Asn
                   150
                                        155
Glu Leu Gln Ser Ala Phe His Leu Pro Glu Gln Asp Leu Ile Tyr Gly
               165
                                   170
Val Phe Thr Thr Asn Val Asn Ser Ile Ala Ala Ser Ala Val Cys Ala
                                185
Phe Asn Leu Ser Ala Ile Ser Gln Ala Phe Asn Gly Pro Phe Arg Tyr
        195
                            200
Gln Glu Asn Pro Arg Ala Ala Trp Leu Pro Ile Ala Asn Pro Ile Pro
                        215
Asn Phe Gln Cys Gly Thr Leu Pro Glu Thr Gly Pro Asn Glu Asn Leu
                    230
                                        235
Thr Glu Arg Ser Leu Gln Asp Ala Gln Arg Leu Phe Leu Met Ser Glu
               245
                                   250
Ala Val Gln Pro Val Thr Pro Glu Pro Cys Val Thr Gln Asp Ser Val
                               265
Arg Phe Ser His Leu Val Val Asp Leu Val Gln Ala Lys Asp Thr Leu
                           280
                                                285
Tyr His Val Leu Tyr Ile Gly Thr Glu Ser Gly Thr Ile Leu Lys Ala
                       295
                                           300
Leu Ser Thr Ala Ser Arg Ser Leu His Gly Cys Tyr Leu Glu Glu Leu
                   310
                                       315
His Val Leu Pro Pro Gly Arg Arg Glu Pro Leu Arg Ser Leu Arg Ile
                325
                                    330
Leu His Ser Ala Arg Ala Leu Phe Val Gly Leu Arg Asp Gly Val Leu
            340
                                345
Arg Val Pro Leu Glu Arg Cys Ala Ala Tyr Arg Ser Gln Gly Ala Cys
                            360
Leu Gly Ala Arg Asp Pro Tyr Cys Gly Trp Asp Gly Lys Gln Gln Arg
                       375
                                           380
Cys Ser Thr Leu Glu Asp Ser Ser Asn Met Ser Leu Trp Thr Gln Asn
                   390
                                       395
Ile Thr Ala Cys Pro Val Arg Asn Val Thr Arg Asp Gly Gly Phe Gly
               405
                                   410
Pro Trp Ser Pro Trp Gln Pro Cys Glu His Leu Asp Gly Asp Asn Ser
           420
                               425
Gly Ser Cys Leu Cys Arg Ala Arg Ser Cys Asp Ser Pro Arg Pro Arg
                            440
Cys Gly Gly Leu Asp Cys Leu Gly Pro Ala Ile His Ile Ala Asn Cys
                        455
Ser Arg Asn Gly Ala Trp Thr Pro Trp Ser Ser Trp Ala Leu Cys Ser
                   470
                                        475
Thr Ser Cys Gly Ile Gly Phe Gln Val Arg Gln Arg Ser Cys Ser Asn
                                   490
Pro Ala Pro Arg His Gly Gly Arg Ile Cys Val Gly Lys Ser Arg Glu
                                505
Glu Arg Phe Cys Asn Glu Asn Thr Pro Cys Pro Val Pro Ile Phe Trp
                            520
Ala Ser Trp Gly Ser Trp Ser Lys Cys Ser Ser Asn Cys Gly Gly
                       535
                                           540
Met Gln Ser Arg Arg Ala Cys Glu Asn Gly Asn Ser Cys Leu Gly
                   550
                                       555
Cys Gly Val Glu Phe Lys Thr Cys Asn Pro Glu Gly Cys Pro Glu Val
                565
                                   570
```

Arg	Arg	Asn	Thr 580	Pro	Trp	Thr	Pro	Trp 585	Leu	Pro	Va1	Asn	Val 590	Thr	Gln
Gly	Gly	Ala 595	Arg	Gln	Glu	Gln	Arg 600	Phe	Arg	Phe	Thr	Cys 605	Arg	Ala	Pro
Leu	Ala 610		Pro	His	Gly	Leu 615	Gln	Phe	Gly	Arg	Arg 620	Arg	Thr	Glu	Thr
Arg 625		Cys	Pro	Ala	Asp		Ser	Gly	Ser	Cys 635	Asp	Thr	Asp	Ala	Leu 640
	Glu	Asp	Leu	Leu 645		Ser	G1y	Ser	Thr 650		Pro	His	Thr	Val 655	Ser
Gly	Gly	Trp	Ala		Trp	Gly	Pro	Trp 665		Ser	Cys	Ser	Arg 670	Asp	Cys
Glu	Leu	Gly 675	Phe	Arg	Val	Arg	Lys 680	Arg	Thr	Cys	Thr	Asn 685	Pro	Glu	Pro
Arg	Asn 690	Gly	Gly	Leu	Pro	Cys 695	Val	Gly	Asp	Ala	Ala 700	Glu	Tyr	Gln	Asp
Cys 705	Asn	Pro	Gln	Ala	Cys 710	Pro	Val	Arg	Gly	Ala 715	Trp	Ser	Cys	Trp	Thr 720
Ser	Trp	Ser	Pro	Cys 725	Ser	Ala	Ser	Cys	Gly 730	Gly	Gly	His	Tyr	Gln 735	Arg
Thr	Arg	Ser	Cys 740	Thr	Ser	Pro	Ala	Pro 745	Ser	Pro	Gly	Glu	Asp 750	Ile	Cys
Leu	Gly	Leu 755	His	Thr	Glu	Glu	Ala 760	Leu	Cys	Ala	Thr	Gln 765	Ala	Cys	Pro
Glu	Gly 770	Trp	Ser	Pro	Trp	Ser 775	Glu	Trp	Ser	Lys	Cys 780	Thr	Asp	Asp	Gly
Ala 785	Gln	Ser	Arg	Ser	Arg 790	His	Cys	Glu	Glu	Leu 795	Leu	Pro	Gly	Ser	Ser 800
Ala	Cys	Ala	Gly	Asn 805	Ser	Ser	Gln	Ser	Arg 810	Pro	Cys	Pro	Tyr	Ser 815	Glu
Ile	Pro	Gly	Phe 820	Asn	Leu	Ile	His	Leu 825	Val	Ala	Thr	Gly	Ile 830	Ser	Cys
Phe	Leu	Gly 835	Ser	Gly	Leu	Leu	Thr 840	Leu	Ala	Val	Tyr	Leu 845	Ser	Cys	Gln
His	Cys 850	Gln	Arg	Gln	Ser	Gln 855	Glu	Ser	Thr	Leu	Val 860	His	Pro	Ala	Thr
Pro 865	Asn	His	Leu	His	Tyr 870	Lys	Gly	Gly	Gly	Thr 875	Pro	Lys	Asn	Glu	Lys 880
Tyr	Thr	Pro	Met	Glu 885	Phe	Lys	Thr	Leu	Asn 890	Lys	Asn	Asn	Leu	Ile 895	Pro
Asp	Asp	Arg	Ala 900		Phe	Tyr	Pro	Leu 905	Gln	Gln	Thr	Asn	Val 910	Tyr	Thr
Thr	Thr	Tyr 915	Tyr	Pro	Ser	Pro	Leu 920	Asn	Lys	His	Ser	Phe 925	Arg	Pro	Glu
Ala	Ser 930	Pro	Gly	Gln	Arg	Cys 935	Phe	Pro	Asn	Ser					
	<210> 17														
	l> 40 2> DI														
			sapi	ens											

<400> 17

cccgcggtctcctcctcctgctctctcgagcgccggtcgggagctagttggagcggggggttggtgccagagccagctccgccgagccgggcgggtcggcagcgatccagcggtgctgggagcccgagcgcagccgggtggggactgcaccggagcgctgag180

					1 - 1 - 1	0.40
	cgttcctgcg					240
	cgctccgcgc					300
	ggaggtcttc					360
	cttgcaggcc					420
	ctctccagct					480
	agcaagcacc					540
	cctggagccc					600
	ggagccagga					660
	gagtgggcct					720
	gagtgtcaga					780
	accaatgcct					840
	gagaagatca					900
	atctcctccc					960
	gccatctacc					1020
	tggcttaatg					1080
	ctgcgggaga					1140
	gtgtgcaaga					1200
	aaggcccggc					1260
	agtgccttcc					1320
	agcatygcgg					1380
	ggcccatttc					1440
	cccaatttcc					1500
	agcctgcagg					1560
	gagccctgtg					1620
	gctaaagaca					1680
	gcgctgtcca					1740
	cccccgggc					1800
	ttcgtggggc					1860
	agccaggggg					1920
	cgttgcagca					1980
	tgtcctgtgc					2040
	tgtgagcact					2100
	tcccctcgac					2160
	tgctccagga					2220
	ggcatcggct					2280
	cgcatctgcg					2340
	gtgcccatct					2400
	ggcatgcagt					2460
	gagttcaaga					2520
	ccgtggctgc					2580
	acctgccgcg					2640
	acgaggacct					2700
	ctcctgcgca					2760
	ccgtggtcgt					2820
	actaacccgg					2880
	gactgcaacc					2940
	ccatgctcag					3000
	gcaccctccc					3060
	acacaggcct					3120
	ggagcccaga					3180
	ggaaacagca					3240
	tccagcatgg					3300
	ggcatctcct					3360
	cactgccagc					3420
	cactacaagg					3480
accaagacc	ctgaacaaga	acaacttgat	ccctgatgac	agagccaact	tctacccatt	3540

```
gcagcagacc aatgtgtaca cgactactta ctacccaagc cccctgaaca aacacagctt
                                                                       3600
 ccggcccgag gcctcacctg gacaacggtg cttccccaac agctgatacc gccgtcctgg
                                                                       3660
 ggacttgggc ttcttgcctt cataaggcac agagcagatg gagatgggac agtggagcca
                                                                       3720
 gtttggtttt ctccctctgc actaggccaa gaacttgctg ccttgcctgt ggggggtccc
                                                                       3780
 atccggcttc agagagetet ggctggcatt gaccatgggg gaaagggetg gtttcagget
                                                                       3840
 gacatatggc cgcaggtcca gttcagccca ggtctctcat ggttatcttc caacccactg
                                                                       3900
 tcacgctgac actatgctgc catgcctggg ctgtggacct actgggcatt tgaggaattg
                                                                       3960
 gagaatggag atggcaagag ggcaggcttt taagtttggg ttggagacaa cttcctgtgg
                                                                      4020
 cccccacaag ctgagtctgg ccttctccag ctggccccaa aaaaggcctt tgct
                                                                       4074
 <210> 18
 <211> 648
 <212> DNA
 <213> homo sapiens
 <400> 18
 atgtggggga ggctctggcc cctcctcctc agcatcctca cagcaactgc agtcccagga
                                                                        60
 ccctcactgc ggagaccgtc tagagaacta gatgccaccc ctcggatgac cataccctat
                                                                       120
gaagagetet etgggaeeeg geaetteaag ggeeaageee agaactaete aacaetgetg
                                                                       180
ctggaggagg cctcagcaag gctgctggtg ggagcccgag gtgccctgtt ctctctcagt
                                                                       240
gccaacgaca taggagatgg ggctcacaaa gagatccact gggaagcctc cccagagatg
                                                                       300
caaagcaaat gtcatcaaaa agggaaaaac aaccagacgg agtgctttaa ccatgtgcgg
                                                                       360
ttcctgcagc ggctcaattc tacccacctc tatgcatgtg ggactcacgc cttccagccc
                                                                       420
ctctgtgcag ccattgatgc tgaggccttc accttgccaa ccagcttcga ggaggggaag
                                                                       480
gagaagtgtc cttatgaccc agcccgtggc ttcacaggcc tcatcattga tggaggcctc
                                                                       540
tacacageca ctaggtatga attecggage attectgaca tecgeeggag eegecaceca
                                                                       600
cactccctga gaactgagga gacaccaatg cattggctca atggttag
                                                                       648
<210> 19
<211> 215
<212> PRT
<213> homo sapiens
<400> 19
Met Trp Gly Arg Leu Trp Pro Leu Leu Leu Ser Ile Leu Thr Ala Thr
                 5
                                    10
Ala Val Pro Gly Pro Ser Leu Arg Arg Pro Ser Arg Glu Leu Asp Ala
            20
                                25
Thr Pro Arg Met Thr Ile Pro Tyr Glu Glu Leu Ser Gly Thr Arg His
                            40
Phe Lys Gly Gln Ala Gln Asn Tyr Ser Thr Leu Leu Leu Glu Glu Ala
                                            60
Ser Ala Arg Leu Leu Val Gly Ala Arg Gly Ala Leu Phe Ser Leu Ser
                    70
                                        75
Ala Asn Asp Ile Gly Asp Gly Ala His Lys Glu Ile His Trp Glu Ala
                                    90
Ser Pro Glu Met Gln Ser Lys Cys His Gln Lys Gly Lys Asn Asn Gln
                                105
Thr Glu Cys Phe Asn His Val Arg Phe Leu Gln Arg Leu Asn Ser Thr
        115
                            120
                                                125
His Leu Tyr Ala Cys Gly Thr His Ala Phe Gln Pro Leu Cys Ala Ala
                        135
Ile Asp Ala Glu Ala Phe Thr Leu Pro Thr Ser Phe Glu Glu Gly Lys
145
                                        155
Glu Lys Cys Pro Tyr Asp Pro Ala Arg Gly Phe Thr Gly Leu Ile Ile
```

```
165
                                     170
                                                          175
 Asp Gly Gly Leu Tyr Thr Ala Thr Arg Tyr Glu Phe Arg Ser Ile Pro
                                 185
 Asp Ile Arg Arg Ser Arg His Pro His Ser Leu Arg Thr Glu Glu Thr
                             200
                                                  205
 Pro Met His Trp Leu Asn Gly
 <210> 20
 <211> 1491
 <212> DNA
 <213> homo sapiens
 <400> 20
 atgtggggga ggctctggcc cctcctcctc agcatcctca cagcaactgc agtcccagga
                                                                        60
 ccctcactgc ggagaccgtc tagagaacta gatgccaccc ctcggatgac cataccctat
                                                                        120
 gaagagetet etgggaeeeg geaetteaag ggeeaageee agaaetaete aacaetgetg
                                                                        180
ctggaggagg cctcagcaag gctgctggtg ggagcccgag gtgccctgtt ctctctcagt
                                                                        240
 gccaacgaca taggagatgg ggctcacaaa gagatccact gggaagcctc cccagagatg
                                                                        300
caaagcaaat gtcatcaaaa agggaaaaac aaccagacgg agtgctttaa ccatgtgcgg
                                                                       360
ttcctgcagc ggctcaattc tacccacctc tatgcatgtg ggactcacgc cttccagccc
                                                                        420
ctctgtgcag ccattgatgc tgaggccttc accttgccaa ccagcttcga ggaggggaag
                                                                       480
gagaagtgtc cttatgaccc agcccgtggc ttcacaggcc tcatcattga tggaggcctc
                                                                       540
tacacageca ctaggtatga attecggage attectgaca teegeeggag eegecaceca
                                                                       600
cactccctga gaactgagga gacaccaatg cattggctca atgatgcgga gtttgtgttc
                                                                       660
tccgtcctcg tgcgggagag caaggccagt gcagtgggtg atgatgacaa ggtgtactac
                                                                       720
ttcttcacgg agcgtgccac tgaggagggc tctggcagct tcactcagag ccgcagcagt
                                                                       780
caccgtgtgg cccgtgtggc tcgygtctgc aagggagacc tgggagggaa gaagatcctg
                                                                       840
cagaagaagt ggactteett eetgaaagee egteteatet geeacattee aetgtatgag
                                                                       900
acactgcgtg gggtctgcag cctggatgct gaaacctcaa gccgtacaca cttctatgca
                                                                       960
gccttcacgc tgagcacaca gtggaagacc ctggaggcct cagccatctg ccgctatgac
                                                                      1020
ctggcagaga tccaggctgt ctttgcagga ccctatatgg aataccagga tggttcccgg
                                                                      1080
egetggggte getatgaggg tggggtgeet gageeeegge etggetegtg tateaeagat
                                                                      1140
tcattgcgca gccaaggcta caattcatcc caagacttgc catccctggt cctggacttt
                                                                      1200
gtaaagttgc acccactgat ggctcggccc gttgtgccca cacgtggacg gcccctgctg
                                                                      1260
ctcaagcgca acatacgcta cacacactt acagggacac ctgtcaccac gcctgctgga
                                                                      1320
cctacctatg acctgctctt tctgggcaca gctgatggct ggatccacaa ggccgtagtc
                                                                      1380
ctgggctctg ggatgcacat tattgaagag acacaagtgt tcagggagtc ccagtctgtg
                                                                      1440
gaaaatctag tcatctctct attgcaggta gcccttctct gtgaccctta a
                                                                      1491
<210> 21
<211> 496
<212> PRT
<213> homo sapiens
<400> 21
Met Trp Gly Arg Leu Trp Pro Leu Leu Leu Ser Ile Leu Thr Ala Thr
                                    10
Ala Val Pro Gly Pro Ser Leu Arg Arg Pro Ser Arg Glu Leu Asp Ala
                                25
Thr Pro Arg Met Thr Ile Pro Tyr Glu Glu Leu Ser Gly Thr Arg His
                            40
Phe Lys Gly Gln Ala Gln Asn Tyr Ser Thr Leu Leu Glu Glu Ala
                        55
                                            60
Ser Ala Arg Leu Leu Val Gly Ala Arg Gly Ala Leu Phe Ser Leu Ser
```

```
Ala Asn Asp Ile Gly Asp Gly Ala His Lys Glu Ile His Trp Glu Ala
                                    90
Ser Pro Glu Met Gln Ser Lys Cys His Gln Lys Gly Lys Asn Asn Gln
                               105
Thr Glu Cys Phe Asn His Val Arg Phe Leu Gln Arg Leu Asn Ser Thr
                           120
His Leu Tyr Ala Cys Gly Thr His Ala Phe Gln Pro Leu Cys Ala Ala
                       135
                                           140
Ile Asp Ala Glu Ala Phe Thr Leu Pro Thr Ser Phe Glu Glu Gly Lys
                   150
                                       155
Glu Lys Cys Pro Tyr Asp Pro Ala Arg Gly Phe Thr Gly Leu Ile Ile
                165
                                    170
Asp Gly Gly Leu Tyr Thr Ala Thr Arg Tyr Glu Phe Arg Ser Ile Pro
            180
                               185
Asp Ile Arg Arg Ser Arg His Pro His Ser Leu Arg Thr Glu Glu Thr
        195
                          200
Pro Met His Trp Leu Asn Asp Ala Glu Phe Val Phe Ser Val Leu Val
                       215
                                           220
Arg Glu Ser Lys Ala Ser Ala Val Gly Asp Asp Asp Lys Val Tyr Tyr
                   230
                                       235
Phe Phe Thr Glu Arg Ala Thr Glu Glu Gly Ser Gly Ser Phe Thr Gln
                245
                                   250
Ser Arg Ser Ser His Arg Val Ala Arg Val Ala Arg Val Cys Lys Gly
                               265
Asp Leu Gly Gly Lys Lys Ile Leu Gln Lys Lys Trp Thr Ser Phe Leu
                            280
Lys Ala Arg Leu Ile Cys His Ile Pro Leu Tyr Glu Thr Leu Arg Gly
                        295
                                           300
Val Cys Ser Leu Asp Ala Glu Thr Ser Ser Arg Thr His Phe Tyr Ala
                   310
                                       315
Ala Phe Thr Leu Ser Thr Gln Trp Lys Thr Leu Glu Ala Ser Ala Ile
               325
                                   330
Cys Arg Tyr Asp Leu Ala Glu Ile Gln Ala Val Phe Ala Gly Pro Tyr
          340
                               345
Met Glu Tyr Gln Asp Gly Ser Arg Arg Trp Gly Arg Tyr Glu Gly Gly
       355
                           360
                                               365
Val Pro Glu Pro Arg Pro Gly Ser Cys Ile Thr Asp Ser Leu Arg Ser
                       375
                                           380
Gln Gly Tyr Asn Ser Ser Gln Asp Leu Pro Ser Leu Val Leu Asp Phe
                    390
                                       395
Val Lys Leu His Pro Leu Met Ala Arg Pro Val Val Pro Thr Arg Gly
                405
                                   410
Arg Pro Leu Leu Lys Arg Asn Ile Arg Tyr Thr His Leu Thr Gly
           420
                               425
Thr Pro Val Thr Thr Pro Ala Gly Pro Thr Tyr Asp Leu Leu Phe Leu
                           440
                                               445
Gly Thr Ala Asp Gly Trp Ile His Lys Ala Val Val Leu Gly Ser Gly
                       455
                                           460
Met His Ile Ile Glu Glu Thr Gln Val Phe Arg Glu Ser Gln Ser Val
                   470
                                       475
Glu Asn Leu Val Ile Ser Leu Leu Gln Val Ala Leu Leu Cys Asp Pro
               485
                                   490
```

<sup>&</sup>lt;210> 22

<sup>&</sup>lt;211> 2109

<sup>&</sup>lt;212> DNA

```
<400> 22
atgtggggga ggctctggcc cctcctcctc agcatcctca cagcaactgc agtcccagga
                                                                      60
eceteactge ggagacegte tagagaacta gatgecacee eteggatgae catacectat
                                                                     120
gaagagetet etgggaeeeg geaetteaag ggeeaageee agaaetaete aacaetgetg
                                                                     180
ctggaggagg cctcagcaag gctgctggtg ggagcccgag gtgccctgtt ctctctcagt
                                                                     240
gccaacgaca taggagatgg ggctcacaaa gagatccact gggaagcctc cccagagatg
                                                                     300
caaagcaaat gtcatcaaaa agggaaaaac aaccagacgg agtgctttaa ccatgtqcqq
                                                                     360
ttcctgcagc ggctcaattc tacccacctc tatgcatgtg ggactcacgc cttccagccc
                                                                     420
ctctgtgcag ccattgatgc tgaggccttc accttgccaa ccagcttcga ggagggaag
                                                                     480
gagaagtgtc cttatgaccc agcccgtggc ttcacaggcc tcatcattga tggaggcctc
                                                                     540
tacacagcca ctaggtatga attccggagc attcctgaca tccgccggag ccgccaccca
                                                                     600
cactccctga gaactgagga gacaccaatg cattggctca atgatgcgga gtttgtgttc
                                                                     660
tccgtcctcg tgcgggagag caaggccagt gcagtgggtg atgatgacaa ggtgtactac
                                                                     720
ttcttcacgg agcgtgccac tgaggagggc tctggcagct tcactcagag ccgcagcagt
                                                                     780
caccgtgtgg cccgtgtggc tcgygtctgc aagggagacc tgggagggaa gaagatcctg
                                                                     840
cagaagaagt ggacttcctt cctgaaagcc cgtctcatct gccacattcc actgtatgag
                                                                     900
acactgogtg gggtctgcag cctggatgct gaaacctcaa gccgtacaca cttctatgca
                                                                     960
geetteaege tgageaeaea gtggaagaee etggaggeet eageeatetg eegetatgae
                                                                    1020
ctggcagaga tccaggctgt ctttgcagga ccctatatgg aataccagga tggttcccgg
                                                                    1080
cgctggggtc gctatgaggg tggggtgcct gagccccggc ctggctcgtg tatcacagat
                                                                    1140
tcattgcgca gccaaggcta caattcatcc caagacttgc catccctggt cctggacttt
                                                                    1200
gtaaagttgc acccactgat ggctcggccc gttgtgccca cacgtggacg gccctgctg
                                                                    1260
ctcaagcgca acatacgcta cacacctt acagggacac ctgtcaccac gcctgctgga
                                                                    1320
cctacctatg acctgctctt tctgggcaca gctgatggct ggatccacaa ggccqtagtc
                                                                    1380
etgggetetg ggatgeacat tattgaagag acacaagtgt teagggagte eeagtetgtg
                                                                   1440
gaaaatctag tcatctctct attgcagcac agcctctatg tgggggctcc tagcggagtc
                                                                   1500
atccagctac cactetecag etgetecege taccgatect getatgactg catettggee
                                                                   1560
cgagacccct actgtggctg ggaccctggc acccatgcct gcgcagcagc caccaccata
                                                                   1620
gccaacaggt cccagggaag caggacagca ctgatacagg acatagagag aggaaatcga
                                                                   1680
ggctgtgaga gcagcaggga tacaggcagg gctctgcagg tccatatggg ctcaatgtca
                                                                   1740
ccaccctctg catggccctg tgtgctggat ggtcctgaaa ccagacaagt cctctgccag
                                                                   1800
1860
cagtgccccc accctcacct tetectggtg cattettgtt teatecetge ttetggactt
                                                                   1920
ggggtaccct cccaattgcc acatcctatc tggtcctctt ccccagcccc atgtggtgac
                                                                   1980
ctctttgtca agagcttggg aacgggccag cctggggagg taagactgca tcactcccct
                                                                   2040
cetetecett cetgtgtggc cettgtgaat cageeteece acteteettg gteattetea
                                                                   2100
agagtatga
                                                                   2109
```

```
<210> 23
```

## <400> 23

```
      Met
      Trp
      Gly
      Arg
      Leu
      Trp
      Pro
      Leu
      Leu
      Leu
      Ser
      Ile
      Leu
      Thr
      Ala
      Thr

      Ala
      Val
      Pro
      Gly
      Pro
      Ser
      Leu
      Arg
      Pro
      Pro
      Ser
      Arg
      Glu
      Leu
      Asp
      Ala

      Ala
      Pro
      Arg
      Met
      Thr
      Ile
      Pro
      Tyr
      Glu
      Glu
      Leu
      Gly
      Thr
      Arg
      His

      Ala
      Fro
      Ile
      I
```

<sup>&</sup>lt;211> 702

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> homo sapiens

Ala Asn Asp Ile Gly Asp Gly Ala His Lys Glu Ile His Trp Glu Ala 85 90 Ser Pro Glu Met Gln Ser Lys Cys His Gln Lys Gly Lys Asn Asn Gln 105 100 Thr Glu Cys Phe Asn His Val Arg Phe Leu Gln Arg Leu Asn Ser Thr 120 His Leu Tyr Ala Cys Gly Thr His Ala Phe Gln Pro Leu Cys Ala Ala 135 140 Ile Asp Ala Glu Ala Phe Thr Leu Pro Thr Ser Phe Glu Glu Gly Lys 150 155 Glu Lys Cys Pro Tyr Asp Pro Ala Arg Gly Phe Thr Gly Leu Ile Ile 170 165 Asp Gly Gly Leu Tyr Thr Ala Thr Arg Tyr Glu Phe Arg Ser Ile Pro 185 Asp Ile Arg Arg Ser Arg His Pro His Ser Leu Arg Thr Glu Glu Thr 200 Pro Met His Trp Leu Asn Asp Ala Glu Phe Val Phe Ser Val Leu Val 215 Arg Glu Ser Lys Ala Ser Ala Val Gly Asp Asp Asp Lys Val Tyr Tyr 235 230 Phe Phe Thr Glu Arg Ala Thr Glu Glu Gly Ser Gly Ser Phe Thr Gln 250 245 Ser Arg Ser Ser His Arg Val Ala Arg Val Ala Arg Val Cys Lys Gly 265 260 Asp Leu Gly Gly Lys Lys Ile Leu Gln Lys Lys Trp Thr Ser Phe Leu 280 Lys Ala Arg Leu Ile Cys His Ile Pro Leu Tyr Glu Thr Leu Arg Gly 295 300 Val Cys Ser Leu Asp Ala Glu Thr Ser Ser Arg Thr His Phe Tyr Ala 310 315 Ala Phe Thr Leu Ser Thr Gln Trp Lys Thr Leu Glu Ala Ser Ala Ile 330 325 Cys Arg Tyr Asp Leu Ala Glu Ile Gln Ala Val Phe Ala Gly Pro Tyr 345 Met Glu Tyr Gln Asp Gly Ser Arg Arg Trp Gly Arg Tyr Glu Gly Gly 360 Val Pro Glu Pro Arg Pro Gly Ser Cys Ile Thr Asp Ser Leu Arg Ser 380 375 Gln Gly Tyr Asn Ser Ser Gln Asp Leu Pro Ser Leu Val Leu Asp Phe 395 390 Val Lys Leu His Pro Leu Met Ala Arg Pro Val Val Pro Thr Arg Gly 405 410 Arg Pro Leu Leu Lys Arg Asn Ile Arg Tyr Thr His Leu Thr Gly 425 Thr Pro Val Thr Thr Pro Ala Gly Pro Thr Tyr Asp Leu Leu Phe Leu 440 Gly Thr Ala Asp Gly Trp Ile His Lys Ala Val Val Leu Gly Ser Gly 455 460 Met His Ile Ile Glu Glu Thr Gln Val Phe Arg Glu Ser Gln Ser Val 475 470 Glu Asn Leu Val Ile Ser Leu Leu Gln His Ser Leu Tyr Val Gly Ala 490 Pro Ser Gly Val Ile Gln Leu Pro Leu Ser Ser Cys Ser Arg Tyr Arg 505 Ser Cys Tyr Asp Cys Ile Leu Ala Arg Asp Pro Tyr Cys Gly Trp Asp 515 520

```
Pro Gly Thr His Ala Cys Ala Ala Ala Thr Thr Ile Ala Asn Arg Ser
                        535
                                             540
Gln Gly Ser Arg Thr Ala Leu Ile Gln Asp Ile Glu Arg Gly Asn Arg
                    550
                                         555
Gly Cys Glu Ser Ser Arg Asp Thr Gly Arg Ala Leu Gln Val His Met
                565
                                     570
Gly Ser Met Ser Pro Pro Ser Ala Trp Pro Cys Val Leu Asp Gly Pro
                                                     590
            580
                                585
Glu Thr Arg Gln Val Leu Cys Gln Pro Pro Lys Pro Cys Val His Ser
                            600
                                                 605
His Ala His Met Glu Glu Cys Leu Ser Ala Gly Leu Gln Cys Pro His
                        615
                                             620
Pro His Leu Leu Val His Ser Cys Phe Ile Pro Ala Ser Gly Leu
625
                    630
                                         635
                                                             640
Gly Val Pro Ser Gln Leu Pro His Pro Ile Trp Ser Ser Pro Ala
                645
                                     650
Pro Cys Gly Asp Leu Phe Val Lys Ser Leu Gly Thr Gly Gln Pro Gly
            660
                                665
Glu Val Arg Leu His His Ser Pro Pro Leu Pro Ser Cys Val Ala Leu
                            680
Val Asn Gln Pro Pro His Ser Pro Trp Ser Phe Ser Arg Val
    690
                        695
<210> 24
<211> 2094
<212> DNA
```

<213> homo sapiens

## <400> 24

60 atgtggggga ggctctggcc cctcctcctc agcatcctca cagcaactgc agtcccagga 120 ccctcactgc ggagaccgtc tagagaacta gatgccaccc ctcggatgac cataccctat 180 gaagagetet etgggaceeg geactteaag ggecaageee agaactaete aacaetgetg 240 ctggaggagg cctcagcaag gctgctggtg ggagcccgag gtgccctgtt ctctctcagt 300 gccaacgaca taggagatgg ggctcacaaa gagatccact gggaagcctc cccagagatg 360 caaagcaaat gtcatcaaaa agggaaaaac aaccagacgg agtgctttaa ccatgtgcgg 420 ttcctgcagc ggctcaattc tacccacctc tatgcatgtg ggactcacgc cttccagccc 480 ctctgtgcag ccattgatgc tgaggccttc accttgccaa ccagcttcga ggaggggaag 540 gagaagtgtc cttatgaccc agcccgtggc ttcacaggcc tcatcattga tggaggcctc tacacaqcca ctaqqtatqa attccqqaqc attcctqaca tccqccqqaq ccgccaccca 600 cactecetga gaactgagga gacaccaatg cattggetca atgatgegga gtttgtgtte 660 720 tccgtcctcg tgcgggagag caaggccagt gcagtgggtg atgatgacaa ggtgtactac 780 ttcttcacgg agcgtgccac tgaggagggc tctggcagct tcactcagag ccgcagcagt 840 caccgtgtgg cccgtgtggc tcgygtctgc aagggagacc tgggagggaa gaagatcctg cagaagaagt ggacttcctt cctgaaagcc cgtctcatct gccacattcc actgtatgag 900 960 acactgcgtg gggtctgcag cctggatgct gaaacctcaa gccgtacaca cttctatgca 1020 gccttcacgc tgagcacaca gtggaagacc ctggaggcct cagccatctg ccgctatgac 1080 ctggcagaga tccaggctgt ctttgcagga ccctatatgg aataccagga tggttcccgg 1140 cgctggggtc gctatgaggg tggggtgcct gagccccggc ctggctcgtg tatcacagat 1200 tcattgcgca gccaaggcta caattcatcc caagacttgc catccctggt cctggacttt 1260 gtaaagttgc acccactgat ggctcggccc gttgtgccca cacgtggacg gcccctgctg ctcaagcgca acatacgcta cacacacctt acagggacac ctgtcaccac gcctgctgga 1320 1380 cctacctatg acctgctctt tctgggcaca gctgatggct ggatccacaa ggccgtagtc ctgggctctg ggatgcacat tattgaagag acacaagtgt tcagggagtc ccagtctgtg 1440 gaaaatctag tcatctctct attgcagcac agcctctatg tgggggctcc tagcggagtc 1500 atccagctac cactetecag etgetecege taccgatect getatgactg catettggee 1560 1620 cgagacccct actgtggctg ggaccctggc acccatgcct gcgcagcagc caccaccata

```
gccaacagga cagcactgat acaggacata gagagaggaa atcgaggctg tgagagcagc
                                                                      1680
agggatacag gcagggctct gcaggtccat atgggctcaa tgtcaccacc ctctgcatgg
                                                                      1740
ccctgtgtgc tggatggtcc tgaaaccaga caagtcctct gccagccacc taagccctgc
                                                                      1800
                                                                      1860
gtacattcac atgcacacat ggaagaatgt ttatcggctg ggctgcagtg cccccaccct
caccttctcc tggtgcattc ttgtttcatc cctgcttctg gacttggggt accctcccaa
                                                                      1920
ttgccacatc ctatctggtc ctcttcccca gccccatgtg gtgacctctt tgtcaagagc
                                                                      1980
ttgggaacgg gccagcctgg ggaggtaaga ctgcatcact cccctcctct cccttcctgt
                                                                      2040
                                                                      2094
qtqqccttq tqaatcaqcc tccccactct ccttggtcat tctcaagagt atga
```

<211> 697

<212> PRT

260

<213> homo sapiens <400> 25 Met Trp Gly Arg Leu Trp Pro Leu Leu Leu Ser Ile Leu Thr Ala Thr 10 Ala Val Pro Gly Pro Ser Leu Arg Arg Pro Ser Arg Glu Leu Asp Ala 20 25 Thr Pro Arg Met Thr Ile Pro Tyr Glu Glu Leu Ser Gly Thr Arg His 45 40 Phe Lys Gly Gln Ala Gln Asn Tyr Ser Thr Leu Leu Glu Glu Ala 55 Ser Ala Arg Leu Leu Val Gly Ala Arg Gly Ala Leu Phe Ser Leu Ser 75 Ala Asn Asp Ile Gly Asp Gly Ala His Lys Glu Ile His Trp Glu Ala 85 Ser Pro Glu Met Gln Ser Lys Cys His Gln Lys Gly Lys Asn Asn Gln 110 105 Thr Glu Cys Phe Asn His Val Arg Phe Leu Gln Arg Leu Asn Ser Thr 120 His Leu Tyr Ala Cys Gly Thr His Ala Phe Gln Pro Leu Cys Ala Ala 135 140 Ile Asp Ala Glu Ala Phe Thr Leu Pro Thr Ser Phe Glu Glu Gly Lys 155 150 Glu Lys Cys Pro Tyr Asp Pro Ala Arg Gly Phe Thr Gly Leu Ile Ile 170 165 Asp Gly Gly Leu Tyr Thr Ala Thr Arg Tyr Glu Phe Arg Ser Ile Pro 190 185 180 Asp Ile Arg Arg Ser Arg His Pro His Ser Leu Arg Thr Glu Glu Thr 205 200 195 Pro Met His Trp Leu Asn Asp Ala Glu Phe Val Phe Ser Val Leu Val 220 215 Arg Glu Ser Lys Ala Ser Ala Val Gly Asp Asp Asp Lys Val Tyr Tyr 235 230 Phe Phe Thr Glu Arg Ala Thr Glu Glu Gly Ser Gly Ser Phe Thr Gln 245 250 Ser Arg Ser Ser His Arg Val Ala Arg Val Ala Arg Val Cys Lys Gly

315

265 Asp Leu Gly Gly Lys Lys Ile Leu Gln Lys Lys Trp Thr Ser Phe Leu

280 Lys Ala Arg Leu Ile Cys His Ile Pro Leu Tyr Glu Thr Leu Arg Gly

Val Cys Ser Leu Asp Ala Glu Thr Ser Ser Arg Thr His Phe Tyr Ala

Ala Phe Thr Leu Ser Thr Gln Trp Lys Thr Leu Glu Ala Ser Ala Ile

295

310

270

				325					330					335	
Cys	Arg	Tyr	Asp 340	Leu	Ala	Glu	Ile	Gln 345	Ala	Val	Phe	Ala	Gly 350	Pro	Tyr
Met	Glu	Tyr 355	Gln	Asp	Gly	Ser	Arg 360	Arg	Trp	Gly	Arg	Tyr 365	Glu	Gly	Gly
Val	Pro 370	Glu	Pro	Arg	Pro	Gly 375	Ser	Cys	Ile	Thr	Asp 380	Ser	Leu	Arg	Ser
Gln 385	Gly	Tyr	Asn	Ser	Ser 390	Gln	Asp	Leu	Pro	Ser 395	Leu	Val	Leu	Asp	Phe 400
Val	Lys	Leu	His	Pro 405	Leu	Met	Ala	Arg	Pro 410	Val	Val	Pro	Thr	Arg 415	Gly
Arg	Pro	Leu	Leu 420	Leu	Lys	Arg	Asn	Ile 425	Arg	Tyr	Thr	His	Leu 430	Thr	Gly
Thr	Pro	Val 435	Thr	Thr	Pro	Ala	Gly 440	Pro	Thr	Tyr	Asp	Leu 445	Leu	Phe	Leu
Gly	Thr 450	Ala	Asp	Gly	Trp	Ile 455	His	Lys	Ala	Val	Val 460	Leu	Gly	Ser	Gly
Met 465	His	Ile	Ile	Glu	Glu 470	Thr	Gln	Val	Phe	Arg 475	Glu	Ser	Gln	Ser	Val 480
Glu	Asn	Leu	Val	Ile 485	Ser	Leu	Leu	Gln	His 490	Ser	Leu	Tyr	Val	Gly 495	Ala
Pro	Ser	Gly	Val 500	Ile	Gln	Leu	Pro	Leu 505	Ser	Ser	Cys	Ser	Arg 510	Tyr	Arg
Ser	Cys	Tyr 515	Asp	Cys	Ile	Leu	Ala 520	Arg	Asp	Pro	Tyr	Cys 525	Gly	Trp	Asp
Pro	Gly 530	Thr	His	Ala	Cys	Ala 535	Ala	Ala	Thr	Thr	Ile 540	Ala	Asn	Arg	Thr
Ala 545	Leu	Ile	Gln	Asp	Ile 550	Glu	Arg	Gly	Asn	Arg 555	Gly	Cys	Glu	Ser	Ser 560
Arg	Asp	Thr	Gly	Arg 565	Ala	Leu	Gln	Val	His 570	Met	Gly	Ser	Met	Ser 575	Pro
Pro	Ser	Ala	Trp 580	Pro	Cys	Val	Leu	Asp 585	Gly	Pro	Glu	Thr	Arg 590	Gln	Va1
Leu	Cys	Gln 595	Pro	Pro	Lys	Pro	Суs 600	Val	His	Ser	His	Ala 605	His	Met	Glu
Glu	Cys 610	Leu	Ser	Ala	Gly	Leu 615	Gln	Cys	Pro	His	Pro 620	His	Leu	Leu	Leu
Val 625	His	Ser	Cys	Phe	I1e 630	Pro	Ala	Ser	Gly	Leu 635	Gly	Val	Pro	Ser	Gln 640
Leu	Pro	His	Pro	Ile 645	Trp	Ser	Ser	Ser	Pro 650	Ala	Pro	Cys	Gly	Asp 655	Leu
Phe	Val	Lys	Ser 660	Leu	Gly	Thr	Gly	Gln 665	Pro	Gly	Glu	Val	Arg 670	Leu	His
His	Ser	Pro 675	Pro	Leu	Pro	Ser	Cys 680	Val	Ala	Leu	Val	Asn 685	Gln	Pro	Pro
His	Ser 690	Pro	Trp	Ser	Phe	Ser 695	Arg	Val							
	0> 20 L> 25														

<212> DNA

<213> homo sapiens

<400> 26

atgtggggga ggctctggcc cetectecte ageatectea eageaactge agteceagga eecteactge ggagacegte tagagaacta gatgecaeee eteggatgae cataceetat

60

```
gaagagctct ctgggacccg gcacttcaag ggccaagccc agaactactc aacactgctg
                                                                       180
ctggaggagg cctcagcaag gctgctggtg ggagcccgag gtgccctgtt ctctctcagt
                                                                       240
gccaacgaca taggagatgg ggctcacaaa gagatccact gggaagcctc cccagagatg
                                                                       300
caaagcaaat gtcatcaaaa agggaaaaac aaccagacgg agtgctttaa ccatgtgcgg
                                                                       360
ttcctgcagc ggctcaattc tacccacctc tatgcatgtg ggactcacgc cttccagccc
                                                                       420
ctctgtgcag ccattgatgc tgaggccttc accttgccaa ccagcttcga ggaggggaag
                                                                       480
gagaagtgtc cttatgaccc agcccgtggc ttcacaggcc tcatcattga tggaggcctc
                                                                       540
tacacageca ctaggtatga attecggage attectgaca tecgeeggag eegecaceca
                                                                       600
cactccctga gaactgagga gacaccaatg cattggctca atgatgcgga gtttgtgttc
                                                                       660
teegteeteg tgegggagag caaggeeagt geagtgggtg atgatgaeaa ggtgtaetae
                                                                       720
ttcttcacgg agcgtgccac tgaggagggc tctggcagct tcactcagag ccgcagcagt
                                                                       780
caccgtgtgg cccgtgtggc tcgygtctgc aagggagacc tgggagggaa gaagatcctg
                                                                       840
cagaagaagt ggacttcctt cctgaaagcc cgtctcatct gccacattcc actgtatgag
                                                                       900
acactgegtg gggtetgeag cetggatget gaaaceteaa geegtacaca ettetatgea
                                                                       960
gccttcacgc tgagcacaca gtggaagacc ctggaggcct cagccatctg ccgctatgac
                                                                      1020
ctggcagaga tccaggctgt ctttgcagga ccctatatgg aataccagga tggttcccgg
                                                                      1080
cgctggggtc gctatgaggg tggggtgcct gagccccggc ctggctcgtg tatcacagat
                                                                      1140
tcattgcgca gccaaggcta caattcatcc caagacttgc catccctggt cctggacttt
                                                                      1200
gtaaagttgc acccactgat ggctcggccc gttgtgccca cacgtggacg gcccctgctg
                                                                      1260
ctcaagcgca acatacgcta cacacactt acagggacac ctgtcaccac gcctgctgga
                                                                      1320
cctacctatg acctgetett tetgggeaca getgatgget ggatecaeaa ggeegtagte
                                                                      1380
ctgggctctg ggatgcacat tattgaagag acacaagtgt tcagggagtc ccagtctgtg
                                                                      1440
gaaaatctag tcatctctct attgcagcac agcctctatg tgggggctcc tagcggagtc
                                                                      1500
atccagctac cactetecag etgetecege taccgatect getatgactg catettggee
                                                                      1560
cgagacccct actgtggctg ggaccctggc acccatgcct gcgcagcagc caccaccata
                                                                      1620
gccaacaggt cccagggaag caggacagca ctgatacagg acatagagag aggaaatcga
                                                                      1680
ggctgtgaga gcagcaggga tacagggcca ccaccaccac tgaagacccg ctctgtgctc
                                                                      1740
cggggtgatg atgtectect geeetgtgae cagecateca acetggeeeg ggeettgtgg
                                                                      1800
ctactcaatg ggagcatggg cctgagcgat gggcagggtg gctaccgtgt gggcgtggac
                                                                      1860
gggctgctgg ttacagatgc acagcctgag cacagtggca actatggctg ctatgccgag
                                                                      1920
gaaaatggcc tccgcaccct gctggcctcc tatagtctca cagtccggcc agccactcct
                                                                      1980
gececagete caaaageeee tgecacacet ggggcacage tggcacetga tgtgagaetg
                                                                      2040
ctctatgtgc tagccattgc cgcgcttggt ggccyctgcc tcatcctggc ctcctcctc
                                                                      2100
ctctatgtgg cctgtctgcg ggaaggcaga cgagggcgcc gacggaaata ctcactgggt
                                                                      2160
cgggccagcc gggcaggagg atctgcggtg caactgcaga cagtctcagg ccagtgtcct
                                                                      2220
ggagaggaag atgagggtga tgatgagggg gctggggggcc tggagggcag ctgtctccag
                                                                      2280
ateatecetg gggagggage cecagececa ceacececae egececeaee gecacegget
                                                                      2340
gagctgacca atggcttggt ggcactgccc agccggctgc ggaggatgaa tggcaatagc
                                                                      2400
tatgtgcttc tgaggcagag caacaatgga gtaccagcag ggccctgctc cttcgccgag
                                                                      2460
gaactcagcc gcatcctgga aaaaaggaag cacacgcagc tcgtggagca gctagatgag
                                                                      2520
agctctgtct ga
                                                                      2532
```

<210> 27 <211> 843

<212> PRT

<213> homo sapiens

<400> 27

 Met
 Trp
 Gly
 Arg
 Leu
 Trp
 Pro
 Leu
 Leu
 Leu
 Ser
 Ile
 Leu
 Thr
 Ala
 Thr

 Ala
 Val
 Pro
 Gly
 Pro
 Ser
 Leu
 Arg
 Arg
 Pro
 Ser
 Arg
 Glu
 Leu
 Asp
 Ala

 Thr
 Pro
 Arg
 Met
 Thr
 Ile
 Pro
 Tyr
 Glu
 Glu
 Leu
 Ser
 Gly
 Thr
 Arg
 His

 35
 Ile
 Ile

Ser Ala Arg Leu Leu Val Gly Ala Arg Gly Ala Leu Phe Ser Leu Ser Ala Asn Asp Ile Gly Asp Gly Ala His Lys Glu Ile His Trp Glu Ala Ser Pro Glu Met Gln Ser Lys Cys His Gln Lys Gly Lys Asn Asn Gln Thr Glu Cys Phe Asn His Val Arg Phe Leu Gln Arg Leu Asn Ser Thr His Leu Tyr Ala Cys Gly Thr His Ala Phe Gln Pro Leu Cys Ala Ala Ile Asp Ala Glu Ala Phe Thr Leu Pro Thr Ser Phe Glu Glu Gly Lys Glu Lys Cys Pro Tyr Asp Pro Ala Arg Gly Phe Thr Gly Leu Ile Ile Asp Gly Gly Leu Tyr Thr Ala Thr Arg Tyr Glu Phe Arg Ser Ile Pro Asp Ile Arg Arg Ser Arg His Pro His Ser Leu Arg Thr Glu Glu Thr Pro Met His Trp Leu Asn Asp Ala Glu Phe Val Phe Ser Val Leu Val Arg Glu Ser Lys Ala Ser Ala Val Gly Asp Asp Asp Lys Val Tyr Tyr Phe Phe Thr Glu Arg Ala Thr Glu Glu Gly Ser Gly Ser Phe Thr Gln Ser Arg Ser Ser His Arg Val Ala Arg Val Ala Arg Val Cys Lys Gly Asp Leu Gly Gly Lys Lys Ile Leu Gln Lys Lys Trp Thr Ser Phe Leu Lys Ala Arg Leu Ile Cys His Ile Pro Leu Tyr Glu Thr Leu Arg Gly Val Cys Ser Leu Asp Ala Glu Thr Ser Ser Arg Thr His Phe Tyr Ala Ala Phe Thr Leu Ser Thr Gln Trp Lys Thr Leu Glu Ala Ser Ala Ile Cys Arg Tyr Asp Leu Ala Glu Ile Gln Ala Val Phe Ala Gly Pro Tyr Met Glu Tyr Gln Asp Gly Ser Arg Arg Trp Gly Arg Tyr Glu Gly Gly Val Pro Glu Pro Arg Pro Gly Ser Cys Ile Thr Asp Ser Leu Arg Ser Gln Gly Tyr Asn Ser Ser Gln Asp Leu Pro Ser Leu Val Leu Asp Phe Val Lys Leu His Pro Leu Met Ala Arg Pro Val Val Pro Thr Arg Gly Arg Pro Leu Leu Lys Arg Asn Ile Arg Tyr Thr His Leu Thr Gly Thr Pro Val Thr Thr Pro Ala Gly Pro Thr Tyr Asp Leu Leu Phe Leu Gly Thr Ala Asp Gly Trp Ile His Lys Ala Val Val Leu Gly Ser Gly Met His Ile Ile Glu Glu Thr Gln Val Phe Arg Glu Ser Gln Ser Val Glu Asn Leu Val Ile Ser Leu Leu Gln His Ser Leu Tyr Val Gly Ala Pro Ser Gly Val Ile Gln Leu Pro Leu Ser Ser Cys Ser Arg Tyr Arg 

```
Ser Cys Tyr Asp Cys Ile Leu Ala Arg Asp Pro Tyr Cys Gly Trp Asp
                          520
                                             525
Pro Gly Thr His Ala Cys Ala Ala Ala Thr Thr Ile Ala Asn Arg Ser
                      535
                                         540
Gln Gly Ser Arg Thr Ala Leu Ile Gln Asp Ile Glu Arg Gly Asn Arg
               550
                                     555
Gly Cys Glu Ser Ser Arg Asp Thr Gly Pro Pro Pro Pro Leu Lys Thr
                                 570
               565
Arg Ser Val Leu Arg Gly Asp Asp Val Leu Leu Pro Cys Asp Gln Pro
                              585
Ser Asn Leu Ala Arg Ala Leu Trp Leu Leu Asn Gly Ser Met Gly Leu
                          600
                                             605
Ser Asp Gly Gln Gly Tyr Arg Val Gly Val Asp Gly Leu Leu Val
                      615
Thr Asp Ala Gln Pro Glu His Ser Gly Asn Tyr Gly Cys Tyr Ala Glu
                  630
                                     635
Glu Asn Gly Leu Arg Thr Leu Leu Ala Ser Tyr Ser Leu Thr Val Arg
              645
                                 650
Pro Ala Thr Pro Ala Pro Ala Pro Lys Ala Pro Ala Thr Pro Gly Ala
                           665
Gln Leu Ala Pro Asp Val Arg Leu Leu Tyr Val Leu Ala Ile Ala Ala
                          680
                                            685
Leu Gly Gly Leu Cys Leu Ile Leu Ala Ser Ser Leu Leu Tyr Val Ala
                      695
Cys Leu Arg Glu Gly Arg Arg Gly Arg Arg Arg Lys Tyr Ser Leu Gly
                  710
                                     715
Arg Ala Ser Arg Ala Gly Gly Ser Ala Val Gln Leu Gln Thr Val Ser
               725
                                  730
Gly Gln Cys Pro Gly Glu Glu Asp Glu Gly Asp Asp Glu Gly Ala Gly
                              745
                                                750
           740
Gly Leu Glu Gly Ser Cys Leu Gln Ile Ile Pro Gly Glu Gly Ala Pro
                         760
                                             765
Ala Pro Pro Pro Pro Pro Pro Pro Pro Pro Ala Glu Leu Thr Asn
        775
                                         780
Gly Leu Val Ala Leu Pro Ser Arg Leu Arg Arg Met Asn Gly Asn Ser
                  790
                                     795
Tyr Val Leu Leu Arg Gln Ser Asn Asn Gly Val Pro Ala Gly Pro Cys
              805
                   810
Ser Phe Ala Glu Glu Leu Ser Arg Ile Leu Glu Lys Arg Lys His Thr
                              825
           820
Gln Leu Val Glu Gln Leu Asp Glu Ser Ser Val
                          840
<210> 28
<211> 2517
<212> DNA
<213> homo sapiens
<400> 28
atgtggggga ggctctggcc cctcctcctc agcatcctca cagcaactgc agtcccagga
                                                                   60
                                                                  120
ccctcactgc ggagaccgtc tagagaacta gatgccaccc ctcggatgac cataccctat
gaagagetet etgggaeeeg geaetteaag ggeeaageee agaaetaete aacaetgetg
                                                                  180
ctggaggagg cctcagcaag gctgctggtg ggagcccgag gtgccctgtt ctctctcagt
                                                                  240
                                                                  300
gccaacgaca taggagatgg ggctcacaaa gagatccact gggaagcctc cccagagatg
                                                                  360
caaagcaaat gtcatcaaaa agggaaaaac aaccagacgg agtgctttaa ccatgtgcgg
```

420

ttcctgcagc ggctcaattc tacccacctc tatgcatgtg ggactcacgc cttccagecc

```
ctctgtgcag ccattgatgc tgaggccttc accttgccaa ccagcttcga ggaggggaag
                                                                      480
gagaagtgtc cttatgaccc agcccgtggc ttcacaggcc tcatcattga tggaggcctc
                                                                      540
tacacagoca ctaggtatga attocggago attoctgaca tocgcoggag cogocacoca
                                                                      600
cactccctga gaactgagga gacaccaatg cattggctca atgatgcgga gtttgtgttc
                                                                      660
                                                                      720
tccgtcctcg tgcgggagag caaggccagt gcagtgggtg atgatgacaa ggtgtactac
ttcttcacgg agcgtgccac tgaggagggc tctggcagct tcactcagag ccgcagcagt
                                                                      780
caccgtgtgg cccgtgtggc tcgygtctgc aagggagacc tgggagggaa gaagatcctg
                                                                      840
cagaagaagt ggactteett cetgaaagee egteteatet geeacattee aetgtatgag
                                                                      900
acactgcgtg gggtctgcag cctggatgct gaaacctcaa gccgtacaca cttctatgca
                                                                      960
gccttcacgc tgagcacaca gtggaagacc ctggaggcct cagccatctg ccgctatgac
                                                                     1020
                                                                     1080
ctggcagaga tccaggctgt ctttgcagga ccctatatgg aataccagga tggttcccgg
                                                                     1140
cgctggggtc gctatgaggg tggggtgcct gagccccggc ctggctcgtg tatcacagat
tcattgcgca gccaaggcta caattcatcc caagacttgc catccctggt cctggacttt
                                                                     1200
gtaaagttgc acccactgat ggctcggccc gttgtgccca cacgtggacg gcccctgctg
                                                                     1260
ctcaagcgca acatacgcta cacacacctt acagggacac ctgtcaccac gcctgctgga
                                                                     1320
cctacctatg acctgctctt tctgggcaca gctgatggct ggatccacaa ggccgtagtc
                                                                     1380
ctgggctctg ggatgcacat tattgaagag acacaagtgt tcagggagtc ccagtctgtg
                                                                      1440
gaaaatctag tcatctctct attgcagcac agcctctatg tgggggctcc tagcggagtc
                                                                      1500
                                                                     1560
atccagctac cactetecag etgetecege taccgatect getatgactg catettggee
cgagacccct actgtggctg ggaccctggc acccatgcct gcgcagcagc caccaccata
                                                                     1620
gccaacagga cagcactgat acaggacata gagagaggaa atcgaggctg tgagagcagc
                                                                      1680
agggatacag ggccaccacc accactgaag acccgctctg tgctccgggg tgatgatgtc
                                                                      1740
ctcctgccct gtgaccagcc atccaacctg gcccgggcct tgtggctact caatgggagc
                                                                      1800
atgggcctga gcgatggca gggtggctac cgtgtgggcg tggacgggct gctggttaca
                                                                      1860
gatgcacagc ctgagcacag tggcaactat ggctgctatg ccgaggaaaa tggcctccgc
                                                                      1920
                                                                      1980
accetgetgg cetectatag teteacagte eggecageca etectgeece agetecaaaa
gcccctgcca cacctggggc acagctggca cctgatgtga gactgctcta tgtgctagcc
                                                                      2040
                                                                      2100
attgccgcgc ttggtggccy ctgcctcatc ctggcctcct ccctcctcta tgtggcctgt
ctgcgggaag gcagacgagg gcgccgacgg aaatactcac tgggtcgggc cagccgggca
                                                                      2160
                                                                      2220
ggaggatetg eggtgeaact geagacagte teaggeeagt gteetggaga ggaagatgag
                                                                      2280
ggtgatgatg agggggctgg gggcctggag ggcagctgtc tccagatcat ccctggggag
ggagccccag ccccaccacc cccaccgccc ccaccgccac cggctgagct gaccaatggc
                                                                      2340
                                                                      2400
ttggtggcac tgcccagccg gctgcggagg atgaatggca atagctatgt gcttctgagg
cagagcaaca atggagtacc agcagggccc tgctccttcg ccgaggaact cagccgcatc
                                                                      2460
                                                                      2517
ctggaaaaaa ggaagcacac gcagctcgtg gagcagctag atgagagctc tgtctga
```

<211> 838

<212> PRT

<213> homo sapiens

100

<400> 29

```
Met Trp Gly Arg Leu Trp Pro Leu Leu Leu Ser Ile Leu Thr Ala Thr
                                    10
                 5
Ala Val Pro Gly Pro Ser Leu Arg Arg Pro Ser Arg Glu Leu Asp Ala
                                                     30
                                25
Thr Pro Arg Met Thr Ile Pro Tyr Glu Glu Leu Ser Gly Thr Arg His
                                                 45
                            40
Phe Lys Gly Gln Ala Gln Asn Tyr Ser Thr Leu Leu Leu Glu Glu Ala
                                             60
                        55
Ser Ala Arg Leu Leu Val Gly Ala Arg Gly Ala Leu Phe Ser Leu Ser
                                        75
                    70
Ala Asn Asp Ile Gly Asp Gly Ala His Lys Glu Ile His Trp Glu Ala
                                     90
Ser Pro Glu Met Gln Ser Lys Cys His Gln Lys Gly Lys Asn Asn Gln
```

```
Thr Glu Cys Phe Asn His Val Arg Phe Leu Gln Arg Leu Asn Ser Thr
                           120
His Leu Tyr Ala Cys Gly Thr His Ala Phe Gln Pro Leu Cys Ala Ala
                      135
                                         140
Ile Asp Ala Glu Ala Phe Thr Leu Pro Thr Ser Phe Glu Glu Gly Lys
                  150
                                      155
Glu Lys Cys Pro Tyr Asp Pro Ala Arg Gly Phe Thr Gly Leu Ile Ile
                                  170
Asp Gly Gly Leu Tyr Thr Ala Thr Arg Tyr Glu Phe Arg Ser Ile Pro
           180
                               185
Asp Ile Arg Arg Ser Arg His Pro His Ser Leu Arg Thr Glu Glu Thr
                           200
Pro Met His Trp Leu Asn Asp Ala Glu Phe Val Phe Ser Val Leu Val
                      215
Arg Glu Ser Lys Ala Ser Ala Val Gly Asp Asp Asp Lys Val Tyr
                                       235
                   230
Phe Phe Thr Glu Arg Ala Thr Glu Glu Gly Ser Gly Ser Phe Thr Gln
                                   250
               245
Ser Arg Ser Ser His Arg Val Ala Arg Val Ala Arg Val Cys Lys Gly
                               265
           260
Asp Leu Gly Gly Lys Lys Ile Leu Gln Lys Lys Trp Thr Ser Phe Leu
                           280
Lys Ala Arg Leu Ile Cys His Ile Pro Leu Tyr Glu Thr Leu Arg Gly
                       295
                                           300
Val Cys Ser Leu Asp Ala Glu Thr Ser Ser Arg Thr His Phe Tyr Ala
                   310
                                       315
Ala Phe Thr Leu Ser Thr Gln Trp Lys Thr Leu Glu Ala Ser Ala Ile
                                   330
               325
Cys Arg Tyr Asp Leu Ala Glu Ile Gln Ala Val Phe Ala Gly Pro Tyr
                               345
           340
Met Glu Tyr Gln Asp Gly Ser Arg Arg Trp Gly Arg Tyr Glu Gly Gly
                          360
                                              365
Val Pro Glu Pro Arg Pro Gly Ser Cys Ile Thr Asp Ser Leu Arg Ser
                      375
                                        380
Gln Gly Tyr Asn Ser Ser Gln Asp Leu Pro Ser Leu Val Leu Asp Phe
                   390
                                       395
Val Lys Leu His Pro Leu Met Ala Arg Pro Val Val Pro Thr Arg Gly
                                    410
Arg Pro Leu Leu Lys Arg Asn Ile Arg Tyr Thr His Leu Thr Gly
            420
                                425
Thr Pro Val Thr Thr Pro Ala Gly Pro Thr Tyr Asp Leu Leu Phe Leu
                           440
Gly Thr Ala Asp Gly Trp Ile His Lys Ala Val Val Leu Gly Ser Gly
                                           460
                       455
Met His Ile Ile Glu Glu Thr Gln Val Phe Arg Glu Ser Gln Ser Val
                   470
                                       475
Glu Asn Leu Val Ile Ser Leu Leu Gln His Ser Leu Tyr Val Gly Ala
               485
                                   490
Pro Ser Gly Val Ile Gln Leu Pro Leu Ser Ser Cys Ser Arg Tyr Arg
                               505
Ser Cys Tyr Asp Cys Ile Leu Ala Arg Asp Pro Tyr Cys Gly Trp Asp
        515
                            520
Pro Gly Thr His Ala Cys Ala Ala Ala Thr Thr Ile Ala Asn Arg Thr
                        535
                                           540
Ala Leu Ile Gln Asp Ile Glu Arg Gly Asn Arg Gly Cys Glu Ser Ser
                    550
```

```
Arg Asp Thr Gly Pro Pro Pro Leu Lys Thr Arg Ser Val Leu Arg
                565
                                    570
Gly Asp Asp Val Leu Leu Pro Cys Asp Gln Pro Ser Asn Leu Ala Arg
            580
                                585
Ala Leu Trp Leu Leu Asn Gly Ser Met Gly Leu Ser Asp Gly Gln Gly
                            600
                                                605
Gly Tyr Arg Val Gly Val Asp Gly Leu Leu Val Thr Asp Ala Gln Pro
                       615
                                            620
Glu His Ser Gly Asn Tyr Gly Cys Tyr Ala Glu Glu Asn Gly Leu Arg
                    630
                                        635
625
Thr Leu Leu Ala Ser Tyr Ser Leu Thr Val Arg Pro Ala Thr Pro Ala
                645
                                    650
Pro Ala Pro Lys Ala Pro Ala Thr Pro Gly Ala Gln Leu Ala Pro Asp
                                665
            660
Val Arg Leu Leu Tyr Val Leu Ala Ile Ala Ala Leu Gly Gly Leu Cys
                            680
                                                685
Leu Ile Leu Ala Ser Ser Leu Leu Tyr Val Ala Cys Leu Arg Glu Gly
                                            700
                        695
Arg Arg Gly Arg Arg Lys Tyr Ser Leu Gly Arg Ala Ser Arg Ala
                    710
                                        715
Gly Gly Ser Ala Val Gln Leu Gln Thr Val Ser Gly Gln Cys Pro Gly
                                    730
                725
Glu Glu Asp Glu Gly Asp Glu Gly Ala Gly Gly Leu Glu Gly Ser
            740
                                745
Cys Leu Gln Ile Ile Pro Gly Glu Gly Ala Pro Ala Pro Pro Pro Pro
                                                765
                            760
Pro Pro Pro Pro Pro Ala Glu Leu Thr Asn Gly Leu Val Ala Leu
                        775
                                            780
Pro Ser Arg Leu Arg Met Asn Gly Asn Ser Tyr Val Leu Leu Arg
                                        795
                    790
Gln Ser Asn Asn Gly Val Pro Ala Gly Pro Cys Ser Phe Ala Glu Glu
                                    810
                805
Leu Ser Arg Ile Leu Glu Lys Arg Lys His Thr Gln Leu Val Glu Gln
            820
                                825
                                                    830
Leu Asp Glu Ser Ser Val
        835
<210> 30
<211> 2613
<212> DNA
<213> homo sapiens
<400> 30
                                                                       60
atgtgggga ggctctggcc cctcctcctc agcatcctca cagcaactgc agtcccagga
                                                                       120
ccctcactgc ggagaccgtc tagagaacta gatgccaccc ctcggatgac cataccctat
gaagagctct ctgggacccg gcacttcaag ggccaagccc agaactactc aacactgctg
                                                                       180
                                                                       240
ctggaggagg cctcagcaag gctgctggtg ggagcccgag gtgccctgtt ctctctcagt
                                                                       300
qccaacqaca taggagatgg ggctcacaaa gagatccact gggaagcctc cccagagatg
caaagcaaat gtcatcaaaa agggaaaaac aaccagacgg agtgctttaa ccatgtgcgg
                                                                       360
                                                                       420
ttcctgcagc ggctcaattc tacccacctc tatgcatgtg ggactcacgc cttccagccc
                                                                       480
ctctgtgcag ccattgatgc tgaggccttc accttgccaa ccagcttcga ggaggggaag
                                                                       540
gagaagtgtc cttatgaccc agcccgtggc ttcacaggcc tcatcattga tggaggcctc
                                                                       600
tacacageca etaggtatga atteeggage atteetgaca teegeeggag eegeeaceca
cactccctga gaactgagga gacaccaatg cattggctca atgatgcgga gtttgtgttc
                                                                       660
                                                                       720
tccgtcctcg tgcgggagag caaggccagt gcagtgggtg atgatgacaa ggtgtactac
```

ttcttcacgg agcgtgccac tgaggagggc tctggcagct tcactcagag ccgcagcagt

```
caccgtgtgg cccgtgtggc tcgygtctgc aagggagacc tgggagggaa gaagatcctg
                                                                       840
cagaagaagt ggactteett eetgaaagee egteteatet geeacattee aetgtatgag
                                                                       900
acactgcgtg gggtctgcag cctggatgct gaaacctcaa gccgtacaca cttctatgca
                                                                       960
gccttcacgc tgagcacaca gtggaagacc ctggaggcct cagccatctg ccgctatgac
                                                                     1020
ctggcagaga tccaggctgt ctttgcagga ccctatatgg aataccagga tggttcccgg
                                                                     1080
cgctggggtc gctatgaggg tggggtgcct gagccccggc ctggctcgtg tatcacagat
                                                                     1140
                                                                     1200
tcattgcgca gccaaggcta caattcatcc caagacttgc catccctggt cctggacttt
gtaaagttgc acccactgat ggctcggccc gttgtgccca cacgtggacg gcccctgctg
                                                                     1260
ctcaagcgca acatacgcta cacacactt acagggacac ctgtcaccac gcctgctgga
                                                                     1320
cctacctatg acctgctctt tctgggcaca gctgatggct ggatccacaa ggccgtagtc
                                                                     1380
ctgggctctg ggatgcacat tattgaagag acacaagtgt tcagggagtc ccagtctgtg
                                                                     1440
gaaaatctag tcatctctct attgcagcac agcctctatg tgggggctcc tagcggagtc
                                                                     1500
atccagctac cactetecag etgetecege taccgatect getatgactg catettggee
                                                                     1560
                                                                     1620
cgagacccct actgtggctg ggaccctggc acccatgcct gcgcagcagc caccaccata
gccaacaggt cccagggaag caggacagca ctgatacagg acatagagag aggaaatcga
                                                                     1680
ggctgtgaga gcagcaggga tacagggcca ccaccaccac tgaagacccg ctctgtgctc
                                                                     1740
cggggtgatg atgtcctcct gccctgtgac cagccatcca acctggcccg ggccttgtgg
                                                                     1800
                                                                     1860
ctactcaatg ggagcatggg cctgagcgat gggcagggtg gctaccgtgt gggcgtggac
gggctgctgg ttacagatgc acagcctgag cacagtggca actatggctg ctatgccgag
                                                                     1920
gaaaatggcc tccgcaccct gctggcctcc tatagtctca cagtccggcc agccactcct
                                                                     1980
                                                                     2040
gccccagctc caaaagcccc tgccacacct ggggcacagc tggcacctga tgtgagactg
ctctatgtgc tagccattgc cgcgcttggt ggccyctgcc tcatcctggc ctcctccctc
                                                                     2100
ctctatgtgg cctgtctgcg ggaaggcaga cgagggcgcc gacggaaata ctcactgggt
                                                                     2160
cgggccagcc gggcaggagg atctgcggtg caactgcaga cagtctcagg cagggctctg
                                                                     2220
                                                                     2280
caggiccata tgggctcaat gicaccaccc tctgcatggc cctgtgtgct ggatggtcct
gaaaccagac aagteetetg ccagecacct aageeetgeg tacatteaca tgcacacatg
                                                                     2340
gaagaatgtt tatcggctgg gctgcagtgc ccccaccctc accttctcct ggtgcattct
                                                                     2400
                                                                     2460
tgtttcatcc ctgcttctgg acttggggta ccctcccaat tgccacatcc tatctggtcc
                                                                     2520
tcttccccag ccccatgtgg tgacctcttt gtcaagagct tgggaacggg ccagcctggg
gaggtaagac tgcatcactc ccctcctct ccttcctgtg tggcccttgt gaatcagcct
                                                                     2580
ccccactctc cttggtcatt ctcaagagta tga
                                                                     2613
```

<210> 31 <211> 870

<212> PRT

<213> homo sapiens

## <400> 31

Met Trp Gly Arg Leu Trp Pro Leu Leu Ser Ile Leu Thr Ala Thr 1 5 10 Ala Val Pro Gly Pro Ser Leu Arg Arg Pro Ser Arg Glu Leu Asp Ala 2.0 25 30 Thr Pro Arg Met Thr Ile Pro Tyr Glu Glu Leu Ser Gly Thr Arg His 40 Phe Lys Gly Gln Ala Gln Asn Tyr Ser Thr Leu Leu Glu Glu Ala 55 60 Ser Ala Arg Leu Leu Val Gly Ala Arg Gly Ala Leu Phe Ser Leu Ser 75 Ala Asn Asp Ile Gly Asp Gly Ala His Lys Glu Ile His Trp Glu Ala 85 90 Ser Pro Glu Met Gln Ser Lys Cys His Gln Lys Gly Lys Asn Asn Gln Thr Glu Cys Phe Asn His Val Arg Phe Leu Gln Arg Leu Asn Ser Thr 120 125 His Leu Tyr Ala Cys Gly Thr His Ala Phe Gln Pro Leu Cys Ala Ala 130 135 140

Ile Asp Ala Glu Ala Phe Thr Leu Pro Thr Ser Phe Glu Glu Gly Lys Glu Lys Cys Pro Tyr Asp Pro Ala Arg Gly Phe Thr Gly Leu Ile Ile Asp Gly Gly Leu Tyr Thr Ala Thr Arg Tyr Glu Phe Arg Ser Ile Pro Asp Ile Arg Arg Ser Arg His Pro His Ser Leu Arg Thr Glu Glu Thr Pro Met His Trp Leu Asn Asp Ala Glu Phe Val Phe Ser Val Leu Val Arg Glu Ser Lys Ala Ser Ala Val Gly Asp Asp Asp Lys Val Tyr Tyr Phe Phe Thr Glu Arg Ala Thr Glu Glu Gly Ser Gly Ser Phe Thr Gln Ser Arg Ser Ser His Arg Val Ala Arg Val Ala Arg Val Cys Lys Gly Asp Leu Gly Gly Lys Lys Ile Leu Gln Lys Lys Trp Thr Ser Phe Leu Lys Ala Arg Leu Ile Cys His Ile Pro Leu Tyr Glu Thr Leu Arg Gly Val Cys Ser Leu Asp Ala Glu Thr Ser Ser Arg Thr His Phe Tyr Ala Ala Phe Thr Leu Ser Thr Gln Trp Lys Thr Leu Glu Ala Ser Ala Ile Cys Arg Tyr Asp Leu Ala Glu Ile Gln Ala Val Phe Ala Gly Pro Tyr Met Glu Tyr Gln Asp Gly Ser Arg Arg Trp Gly Arg Tyr Glu Gly Gly Val Pro Glu Pro Arg Pro Gly Ser Cys Ile Thr Asp Ser Leu Arg Ser Gln Gly Tyr Asn Ser Ser Gln Asp Leu Pro Ser Leu Val Leu Asp Phe Val Lys Leu His Pro Leu Met Ala Arg Pro Val Val Pro Thr Arg Gly Arg Pro Leu Leu Lys Arg Asn Ile Arg Tyr Thr His Leu Thr Gly Thr Pro Val Thr Thr Pro Ala Gly Pro Thr Tyr Asp Leu Leu Phe Leu Gly Thr Ala Asp Gly Trp Ile His Lys Ala Val Val Leu Gly Ser Gly Met His Ile Ile Glu Glu Thr Gln Val Phe Arg Glu Ser Gln Ser Val Glu Asn Leu Val Ile Ser Leu Leu Gln His Ser Leu Tyr Val Gly Ala Pro Ser Gly Val Ile Gln Leu Pro Leu Ser Ser Cys Ser Arg Tyr Arg Ser Cys Tyr Asp Cys Ile Leu Ala Arg Asp Pro Tyr Cys Gly Trp Asp Pro Gly Thr His Ala Cys Ala Ala Ala Thr Thr Ile Ala Asn Arg Ser Gln Gly Ser Arg Thr Ala Leu Ile Gln Asp Ile Glu Arg Gly Asn Arg Gly Cys Glu Ser Ser Arg Asp Thr Gly Pro Pro Pro Leu Lys Thr Arg Ser Val Leu Arg Gly Asp Asp Val Leu Leu Pro Cys Asp Gln Pro 

```
Ser Asn Leu Ala Arg Ala Leu Trp Leu Leu Asn Gly Ser Met Gly Leu
                           600
Ser Asp Gly Gln Gly Gly Tyr Arg Val Gly Val Asp Gly Leu Leu Val
                       615
                                           620
Thr Asp Ala Gln Pro Glu His Ser Gly Asn Tyr Gly Cys Tyr Ala Glu
                   630
                                       635
Glu Asn Gly Leu Arg Thr Leu Leu Ala Ser Tyr Ser Leu Thr Val Arg
                                   650
               645
Pro Ala Thr Pro Ala Pro Ala Pro Lys Ala Pro Ala Thr Pro Gly Ala
                                665
            660
Gln Leu Ala Pro Asp Val Arg Leu Leu Tyr Val Leu Ala Ile Ala Ala
                            680
                                                685
Leu Gly Gly Leu Cys Leu Ile Leu Ala Ser Ser Leu Leu Tyr Val Ala
                                            700
                        695
Cys Leu Arg Glu Gly Arg Arg Gly Arg Arg Lys Tyr Ser Leu Gly
                   710
                                       715
Arg Ala Ser Arg Ala Gly Gly Ser Ala Val Gln Leu Gln Thr Val Ser
                725
                                   730
Gly Arg Ala Leu Gln Val His Met Gly Ser Met Ser Pro Pro Ser Ala
                               745
           740
Trp Pro Cys Val Leu Asp Gly Pro Glu Thr Arg Gln Val Leu Cys Gln
                           760
                                               765
Pro Pro Lys Pro Cys Val His Ser His Ala His Met Glu Glu Cys Leu
                                            780
                        775
Ser Ala Gly Leu Gln Cys Pro His Pro His Leu Leu Val His Ser
                                        795
                    790
Cys Phe Ile Pro Ala Ser Gly Leu Gly Val Pro Ser Gln Leu Pro His
                805
                                    810
Pro Ile Trp Ser Ser Pro Ala Pro Cys Gly Asp Leu Phe Val Lys
                               825
Ser Leu Gly Thr Gly Gln Pro Gly Glu Val Arg Leu His His Ser Pro
                           840
                                                845
Pro Leu Pro Ser Cys Val Ala Leu Val Asn Gln Pro Pro His Ser Pro
                        855
                                            860
Trp Ser Phe Ser Arg Val
865
                    870
<210> 32
<211> 2598
<212> DNA
<213> homo sapiens
<400> 32
                                                                       60
atgtggggga ggctctggcc cctcctcctc agcatcctca cagcaactgc agtcccagga
ccctcactgc ggagaccgtc tagagaacta gatgccaccc ctcggatgac cataccctat
                                                                      120
                                                                      180
gaagagetet etgggaeeeg geaetteaag ggeeaageee agaaetaete aacaetgetg
                                                                      240
ctggaggagg cctcagcaag gctgctggtg ggagcccgag gtgccctgtt ctctctcagt
gccaacgaca taggagatgg ggctcacaaa gagatccact gggaagcctc cccagagatg
                                                                      300
                                                                      360
caaagcaaat gtcatcaaaa agggaaaaac aaccagacgg agtgctttaa ccatgtgcgg
                                                                      420
ttcctgcagc ggctcaattc tacccacctc tatgcatgtg ggactcacgc cttccagccc
                                                                      480
ctctgtgcag ccattgatgc tgaggccttc accttgccaa ccagcttcga ggaggggaag
                                                                      540
gagaaqtgtc cttatgaccc agcccgtggc ttcacaggcc tcatcattga tggaggcctc
                                                                      600
tacacagcca ctaggtatga attccggagc attcctgaca tccgccggag ccgccaccca
                                                                      660
cactccctga gaactgagga gacaccaatg cattggctca atgatgcgga gtttgtgttc
tecgteeteg tgegggagag caaggeeagt geagtgggtg atgatgacaa ggtgtactae
                                                                      720
```

780

ttcttcacgg agcgtgccac tgaggagggc tctggcagct tcactcagag ccgcagcagt

```
840
caccgtgtgg cccgtgtggc tcgygtctgc aagggagacc tgggagggaa gaagatcctg
                                                                       900
cagaagaagt ggacttcctt cctgaaagcc cgtctcatct gccacattcc actgtatgag
                                                                      960
acactgcgtg gggtctgcag cctggatgct gaaacctcaa gccgtacaca cttctatgca
                                                                     1020
gccttcacgc tgagcacaca gtggaagacc ctggaggcct cagccatctg ccgctatgac
ctggcagaga tccaggctgt ctttgcagga ccctatatgg aataccagga tggttcccgg
                                                                     1080
cgctggggtc gctatgaggg tggggtgcct gagccccggc ctggctcgtg tatcacagat
                                                                     1140
tcattgcgca gccaaggcta caattcatcc caagacttgc catccctggt cctggacttt
                                                                     1200
gtaaagttgc acccactgat ggctcggccc gttgtgccca cacgtggacg gcccctgctg
                                                                     1260
ctcaagcgca acatacgcta cacaccctt acagggacac ctgtcaccac gcctgctgga
                                                                     1320
cctacctatg acctgctctt tctgggcaca gctgatggct ggatccacaa ggccgtagtc
                                                                     1380
ctgggctctg ggatgcacat tattgaagag acacaagtgt tcagggagtc ccagtctgtg
                                                                     1440
gaaaatctag tcatctctct attgcagcac agcctctatg tgggggctcc tagcggagtc
                                                                     1500
atccagctac cactetecag etgetecege taccgatect getatgactg catettggee
                                                                     1560
cgagacccct actgtggctg ggaccctggc acccatgcct gcgcagcagc caccaccata
                                                                     1620
                                                                     1680
gccaacagga cagcactgat acaggacata gagagaggaa atcgaggctg tgagagcagc
                                                                     1740
agggatacag ggccaccacc accactgaag accegetetg tgeteegggg tgatgatgte
                                                                     1800
ctcctgccct gtgaccagcc atccaacctg gcccgggcct tgtggctact caatgggagc
                                                                     1860
atgggcctga gcgatgggca gggtggctac cgtgtgggcg tggacgggct gctggttaca
                                                                     1920
gatgcacage ctgagcacag tggcaactat ggctgctatg ccgaggaaaa tggcctccgc
                                                                     1980
accetgetgg cetectatag teteacagte eggeeageea eteetgeece ageteeaaaa
                                                                     2040
gcccctgcca cacctggggc acagctggca cctgatgtga gactgctcta tgtgctagcc
                                                                     2100
attgccgcgc ttggtggccy ctgcctcatc ctggcctcct ccctcctcta tgtggcctgt
                                                                     2160
ctgcgggaag gcagacgagg gcgccgacgg aaatactcac tgggtcgggc cagccgggca
                                                                     2220
ggaggatctg cggtgcaact gcagacagtc tcaggcaggg ctctgcaggt ccatatgggc
                                                                     2280
tcaatgtcac caccctctgc atggccctgt gtgctggatg gtcctgaaac cagacaagtc
                                                                     2340
ctctgccagc cacctaagcc ctgcgtacat tcacatgcac acatggaaga atgtttatcg
getgggetge agtgeeecea ceeteacett eteetggtge attettgttt eateeetget
                                                                     2400
                                                                     2460
totggacttg gggtaccctc ccaattgcca catcctatct ggtcctcttc cccagcccca
                                                                     2520
tgtggtgacc tctttgtcaa gagcttggga acgggccagc ctggggaggt aagactgcat
                                                                     2580
cacteccete etetecette etgtgtggee ettgtgaate ageeteecea eteteettgg
                                                                     2598
tcattctcaa gagtatga
```

<211> 865

<212> PRT

<213> homo sapiens

## <400> 33

Met Trp Gly Arg Leu Trp Pro Leu Leu Ser Ile Leu Thr Ala Thr 1 5 10 Ala Val Pro Gly Pro Ser Leu Arg Arg Pro Ser Arg Glu Leu Asp Ala 25 20 Thr Pro Arg Met Thr Ile Pro Tyr Glu Glu Leu Ser Gly Thr Arg His 40 Phe Lys Gly Gln Ala Gln Asn Tyr Ser Thr Leu Leu Leu Glu Glu Ala 55 60 Ser Ala Arg Leu Leu Val Gly Ala Arg Gly Ala Leu Phe Ser Leu Ser 70 75 Ala Asn Asp Ile Gly Asp Gly Ala His Lys Glu Ile His Trp Glu Ala 90 85 Ser Pro Glu Met Gln Ser Lys Cys His Gln Lys Gly Lys Asn Asn Gln 105 Thr Glu Cys Phe Asn His Val Arg Phe Leu Gln Arg Leu Asn Ser Thr 120 125 His Leu Tyr Ala Cys Gly Thr His Ala Phe Gln Pro Leu Cys Ala Ala 130 135 140

```
Ile Asp Ala Glu Ala Phe Thr Leu Pro Thr Ser Phe Glu Glu Gly Lys
                  150
                                     155
Glu Lys Cys Pro Tyr Asp Pro Ala Arg Gly Phe Thr Gly Leu Ile Ile
              165
                                 170
Asp Gly Gly Leu Tyr Thr Ala Thr Arg Tyr Glu Phe Arg Ser Ile Pro
           180
                             185
Asp Ile Arg Arg Ser Arg His Pro His Ser Leu Arg Thr Glu Glu Thr
                         200
                                   205
Pro Met His Trp Leu Asn Asp Ala Glu Phe Val Phe Ser Val Leu Val
                       215
Arg Glu Ser Lys Ala Ser Ala Val Gly Asp Asp Asp Lys Val Tyr Tyr
                  230
                                      235
Phe Phe Thr Glu Arg Ala Thr Glu Glu Gly Ser Gly Ser Phe Thr Gln
               245
                                  250
Ser Arg Ser Ser His Arg Val Ala Arg Val Ala Arg Val Cys Lys Gly
   260 . 265
Asp Leu Gly Gly Lys Lys Ile Leu Gln Lys Lys Trp Thr Ser Phe Leu
                          280
Lys Ala Arg Leu Ile Cys His Ile Pro Leu Tyr Glu Thr Leu Arg Gly
                     295
                                         300
Val Cys Ser Leu Asp Ala Glu Thr Ser Ser Arg Thr His Phe Tyr Ala
                  310
                                     315
Ala Phe Thr Leu Ser Thr Gln Trp Lys Thr Leu Glu Ala Ser Ala Ile
               325
                                  330
Cys Arg Tyr Asp Leu Ala Glu Ile Gln Ala Val Phe Ala Gly Pro Tyr
           340
                              345
Met Glu Tyr Gln Asp Gly Ser Arg Arg Trp Gly Arg Tyr Glu Gly Gly
                          360
Val Pro Glu Pro Arg Pro Gly Ser Cys Ile Thr Asp Ser Leu Arg Ser
                      375
                                         380
Gln Gly Tyr Asn Ser Ser Gln Asp Leu Pro Ser Leu Val Leu Asp Phe
                                     395
                  390
Val Lys Leu His Pro Leu Met Ala Arg Pro Val Val Pro Thr Arg Gly
               405
                                  410
Arg Pro Leu Leu Lys Arg Asn Ile Arg Tyr Thr His Leu Thr Gly
                             425
          420
Thr Pro Val Thr Thr Pro Ala Gly Pro Thr Tyr Asp Leu Leu Phe Leu
                          440
Gly Thr Ala Asp Gly Trp Ile His Lys Ala Val Val Leu Gly Ser Gly
                      455
Met His Ile Ile Glu Glu Thr Gln Val Phe Arg Glu Ser Gln Ser Val
465
                   470
                                      475
Glu Asn Leu Val Ile Ser Leu Leu Gln His Ser Leu Tyr Val Gly Ala
                                  490
Pro Ser Gly Val Ile Gln Leu Pro Leu Ser Ser Cys Ser Arg Tyr Arg
          500
                              505
Ser Cys Tyr Asp Cys Ile Leu Ala Arg Asp Pro Tyr Cys Gly Trp Asp
                          520
Pro Gly Thr His Ala Cys Ala Ala Ala Thr Thr Ile Ala Asn Arg Thr
                      535
                                          540
Ala Leu Ile Gln Asp Ile Glu Arg Gly Asn Arg Gly Cys Glu Ser Ser
                  550
Arg Asp Thr Gly Pro Pro Pro Leu Lys Thr Arg Ser Val Leu Arg
               565
                                  570
Gly Asp Asp Val Leu Leu Pro Cys Asp Gln Pro Ser Asn Leu Ala Arg
           580
                               585
```

<400> 35

```
Ala Leu Trp Leu Leu Asn Gly Ser Met Gly Leu Ser Asp Gly Gln Gly
                           600
Gly Tyr Arg Val Gly Val Asp Gly Leu Leu Val Thr Asp Ala Gln Pro
                      615
                                          620
Glu His Ser Gly Asn Tyr Gly Cys Tyr Ala Glu Glu Asn Gly Leu Arg
                  630
                                      635
Thr Leu Leu Ala Ser Tyr Ser Leu Thr Val Arg Pro Ala Thr Pro Ala
              645
                                 650
Pro Ala Pro Lys Ala Pro Ala Thr Pro Gly Ala Gln Leu Ala Pro Asp
           660
                 665
Val Arg Leu Leu Tyr Val Leu Ala Ile Ala Ala Leu Gly Gly Leu Cys
       675
                           680
                                              685
Leu Ile Leu Ala Ser Ser Leu Leu Tyr Val Ala Cys Leu Arg Glu Gly
                       695
Arg Arg Gly Arg Arg Lys Tyr Ser Leu Gly Arg Ala Ser Arg Ala
                   710
                                      715
Gly Ser Ala Val Gln Leu Gln Thr Val Ser Gly Arg Ala Leu Gln
               725
                                  730
Val His Met Gly Ser Met Ser Pro Pro Ser Ala Trp Pro Cys Val Leu
           740
                              745
Asp Gly Pro Glu Thr Arg Gln Val Leu Cys Gln Pro Pro Lys Pro Cys
      755
                          760
                                             765
Val His Ser His Ala His Met Glu Glu Cys Leu Ser Ala Gly Leu Gln
                      775
                                          780
Cys Pro His Pro His Leu Leu Val His Ser Cys Phe Ile Pro Ala
                   790
                                      795
Ser Gly Leu Gly Val Pro Ser Gln Leu Pro His Pro Ile Trp Ser Ser
                                  810
Ser Pro Ala Pro Cys Gly Asp Leu Phe Val Lys Ser Leu Gly Thr Gly
                              825
           820
Gln Pro Gly Glu Val Arg Leu His His Ser Pro Pro Leu Pro Ser Cys
                  840
Val Ala Leu Val Asn Gln Pro Pro His Ser Pro Trp Ser Phe Ser Arg
  850
                      855
                                          860
Va1
865
<210> 34
<211> 351
<212> DNA
<213> homo sapiens
<400> 34
                                                                     60
atgcaaagca aatgtcatca aaaagggaaa aacaaccaga cggagtgctt taaccatgtg
cggttcctgc agcggctcaa ttctacccac ctctatgcat gtgggactca cgccttccag
                                                                    120
cccctctgtg cagccattga tgctgaggcc ttcaccttgc caaccagctt cgaggagggg
                                                                    180
                                                                    240
aaggagaagt gtccttatga cccagcccgt ggcttcacag gcctcatcat tgatggaggc
ctctacacag ccactaggta tgaattccgg agcattcctg acatccgccg gagccgccac
                                                                    300
                                                                    351
ccacactccc tgagaactga ggagacacca atgcattggc tcaatggtta g
<210> 35
<211> 116
<212> PRT
<213> homo sapiens
```

35

```
Met Gln Ser Lys Cys His Gln Lys Gly Lys Asn Asn Gln Thr Glu Cys
                                    10
Phe Asn His Val Arg Phe Leu Gln Arg Leu Asn Ser Thr His Leu Tyr
                                25
Ala Cys Gly Thr His Ala Phe Gln Pro Leu Cys Ala Ala Ile Asp Ala
                            40
Glu Ala Phe Thr Leu Pro Thr Ser Phe Glu Glu Gly Lys Glu Lys Cys
                        55
                                            60
Pro Tyr Asp Pro Ala Arg Gly Phe Thr Gly Leu Ile Ile Asp Gly Gly
                    70
                                        75
Leu Tyr Thr Ala Thr Arg Tyr Glu Phe Arg Ser Ile Pro Asp Ile Arg
                                    90
Arg Ser Arg His Pro His Ser Leu Arg Thr Glu Glu Thr Pro Met His
                                105
Trp Leu Asn Gly
        115
<210> 36
<211> 1194
<212> DNA
<213> homo sapiens
<400> 36
                                                                        60
atgcaaagca aatgtcatca aaaagggaaa aacaaccaga cggagtgctt taaccatgtg
cggttcctgc agcggctcaa ttctacccac ctctatgcat gtgggactca cgccttccag
                                                                       120
cccctctgtg cagccattga tgctgaggcc ttcaccttgc caaccagctt cgaggagggg
                                                                       180
aaggagaagt gtccttatga cccagcccgt ggcttcacag gcctcatcat tgatggaggc
                                                                       240
ctctacacag ccactaggta tgaattccgg agcattcctg acatccgccg gagccgccac
                                                                       300
ccacactccc tgagaactga ggagacacca atgcattggc tcaatgatgc ggagtttgtg
                                                                       360
ttctccgtcc tcgtgcggga gagcaaggcc agtgcagtgg gtgatgatga caaggtgtac
                                                                       420
tacttettea eggagegtge caetgaggag ggetetggea getteactea gageegeage
                                                                       480
agtcaccgtg tggcccgtgt ggctcgygtc tgcaagggag acctgggagg gaagaagatc
                                                                       540
ctgcagaaga agtggacttc cttcctgaaa gcccgtctca tctgccacat tccactgtat
                                                                       600
gagacactgc gtggggtctg cagcctggat gctgaaacct caagccgtac acacttctat
                                                                       660
gcagccttca cgctgagcac acagtggaag accctggagg cctcagccat ctgccgctat
                                                                       720
gacctggcag agatccaggc tgtctttgca ggaccctata tggaatacca ggatggttcc
                                                                       780
                                                                       840
cggcgctggg gtcgctatga gggtggggtg cctgagccc ggcctggctc gtgtatcaca
gattcattgc gcagccaagg ctacaattca tcccaagact tgccatccct ggtcctggac
                                                                       900
tttgtaaagt tgcacccact gatggctcgg cccgttgtgc ccacacgtgg acggcccctg
                                                                       960
ctgctcaagc gcaacatacg ctacacaca cttacaggga cacctgtcac cacgcctgct
                                                                      1020
ggacctacct atgacctgct ctttctgggc acagctgatg gctggatcca caaggccgta
                                                                      1080
gtcctgggct ctgggatgca cattattgaa gagacacaag tgttcaggga gtcccagtct
                                                                      1140
gtggaaaatc tagtcatctc tctattgcag gtagcccttc tctgtgaccc ttaa
                                                                      1194
<210> 37
<211> 397
<212> PRT
<213> homo sapiens
<400> 37
Met Gln Ser Lys Cys His Gln Lys Gly Lys Asn Asn Gln Thr Glu Cys
                 5
                                    10
Phe Asn His Val Arg Phe Leu Gln Arg Leu Asn Ser Thr His Leu Tyr
                                25
Ala Cys Gly Thr His Ala Phe Gln Pro Leu Cys Ala Ala Ile Asp Ala
```

```
Glu Ala Phe Thr Leu Pro Thr Ser Phe Glu Glu Gly Lys Glu Lys Cys
                        55
Pro Tyr Asp Pro Ala Arg Gly Phe Thr Gly Leu Ile Ile Asp Gly Gly
                   70
                                       75
Leu Tyr Thr Ala Thr Arg Tyr Glu Phe Arg Ser Ile Pro Asp Ile Arg
               85
                                   90
Arg Ser Arg His Pro His Ser Leu Arg Thr Glu Glu Thr Pro Met His
                               105
           100
Trp Leu Asn Asp Ala Glu Phe Val Phe Ser Val Leu Val Arg Glu Ser
                                              125
                          120
Lys Ala Ser Ala Val Gly Asp Asp Asp Lys Val Tyr Tyr Phe Phe Thr
                        135
                                           140
Glu Arg Ala Thr Glu Glu Gly Ser Gly Ser Phe Thr Gln Ser Arg Ser
                   150
Ser His Arg Val Ala Arg Val Ala Arg Val Cys Lys Gly Asp Leu Gly
                                   170
               165
Gly Lys Lys Ile Leu Gln Lys Lys Trp Thr Ser Phe Leu Lys Ala Arg
                               185
Leu Ile Cys His Ile Pro Leu Tyr Glu Thr Leu Arg Gly Val Cys Ser
                          200
                                              205
Leu Asp Ala Glu Thr Ser Ser Arg Thr His Phe Tyr Ala Ala Phe Thr
                       215
                                           220
Leu Ser Thr Gln Trp Lys Thr Leu Glu Ala Ser Ala Ile Cys Arg Tyr
                                       235
                   230
Asp Leu Ala Glu Ile Gln Ala Val Phe Ala Gly Pro Tyr Met Glu Tyr
                                    250
                245
Gln Asp Gly Ser Arg Arg Trp Gly Arg Tyr Glu Gly Gly Val Pro Glu
                                265
            260
Pro Arg Pro Gly Ser Cys Ile Thr Asp Ser Leu Arg Ser Gln Gly Tyr
                           280
                                                285
Asn Ser Ser Gln Asp Leu Pro Ser Leu Val Leu Asp Phe Val Lys Leu
                       295
His Pro Leu Met Ala Arg Pro Val Val Pro Thr Arg Gly Arg Pro Leu
                   310
                                       315
Leu Leu Lys Arg Asn Ile Arg Tyr Thr His Leu Thr Gly Thr Pro Val
                                   330
               325
Thr Thr Pro Ala Gly Pro Thr Tyr Asp Leu Leu Phe Leu Gly Thr Ala
                               345
Asp Gly Trp Ile His Lys Ala Val Val Leu Gly Ser Gly Met His Ile
                                                365
                            360
Ile Glu Glu Thr Gln Val Phe Arg Glu Ser Gln Ser Val Glu Asn Leu
Val Ile Ser Leu Leu Gln Val Ala Leu Leu Cys Asp Pro
                    390
<210> 38
<211> 1812
<212> DNA
<213> homo sapiens
<400> 38
atgcaaagca aatgtcatca aaaagggaaa aacaaccaga cggagtgctt taaccatgtg
eggtteetge ageggeteaa ttetaeceae etetatgeat gtgggaetea egeetteeag
cccctctgtg cagccattga tgctgaggcc ttcaccttgc caaccagctt cgaggagggg
```

aaggagaagt gtccttatga cccagcccgt ggcttcacag gcctcatcat tgatggaggc

ctctacacag ccactaggta tgaattccgg agcattcctg acatccgccg gagccgccac

60

120 180

240

```
360
ccacactccc tgagaactga ggagacacca atgcattggc tcaatgatgc ggagtttgtg
                                                                       420
ttctccgtcc tcgtgcggga gagcaaggcc agtgcagtgg gtgatgatga caaggtgtac
                                                                       480
tacttettea eggagegtge caetgaggag ggetetggea getteaetea gageegeage
                                                                       540
agtcaccgtg tggcccgtgt ggctcgygtc tgcaagggag acctgggagg gaagaagatc
ctgcagaaga agtggacttc cttcctgaaa gcccgtctca tctgccacat tccactgtat
                                                                       600
gagacactgc gtggggtctg cagcctggat gctgaaacct caagccgtac acacttctat
                                                                       660
gcagcettea egetgageae acagtggaag accetggagg ceteageeat etgeegetat
                                                                       720
                                                                       780
gacctggcag agatccaggc tgtctttgca ggaccctata tggaatacca ggatggttcc
cggcgctggg gtcgctatga gggtggggtg cctgagccc ggcctggctc gtgtatcaca
                                                                       840
gattcattgc gcagccaagg ctacaattca tcccaagact tgccatccct ggtcctggac
                                                                       900
                                                                       960
tttgtaaagt tgcacccact gatggctcgg cccgttgtgc ccacacgtgg acggcccctg
ctgctcaagc gcaacatacg ctacacacac cttacaggga cacctgtcac cacgcctgct
                                                                     1020
                                                                     1080
ggacctacct atgacctgct ctttctgggc acagctgatg gctggatcca caaggccgta
                                                                     1140
gtcctgggct ctgggatgca cattattgaa gagacacaag tgttcaggga gtcccagtct
gtggaaaatc tagtcatctc tctattgcag cacagcctct atgtgggggc tcctagcgga
                                                                     1200
gteatecage taccactete cagetgetee egetacegat cetgetatga etgeatettg
                                                                     1260
gcccgagacc cctactgtgg ctgggaccct ggcacccatg cctgcgcagc agccaccacc
                                                                     1320
atagccaaca ggtcccaggg aagcaggaca gcactgatac aggacataga gagaggaaat
                                                                     1380
cgaggctgtg agagcagcag ggatacaggc agggctctgc aggtccatat gggctcaatg
                                                                     1440
teaceaecet etgeatggee etgtgtgetg gatggteetg aaaceagaea agteetetge
                                                                     1500
                                                                     1560
cagccaccta agccctgcgt acattcacat gcacacatgg aagaatgttt atcggctggg
                                                                     1620
ctgcagtgcc cccaccctca ccttctcctg gtgcattctt gtttcatccc tgcttctgga
cttggggtac cctcccaatt gccacatcct atctggtcct cttccccagc cccatgtggt
                                                                     1680
gacctctttg tcaagagett gggaacgggc cagcctgggg aggtaagact gcatcactcc
                                                                     1740
                                                                     1800
cctcctctcc cttcctgtgt ggcccttgtg aatcagcctc cccactctcc ttggtcattc
tcaagagtat ga
                                                                     1812
```

<211> 603

<212> PRT

<213> homo sapiens

<400> 39

Met Gln Ser Lys Cys His Gln Lys Gly Lys Asn Asn Gln Thr Glu Cys 5 1.0 Phe Asn His Val Arg Phe Leu Gln Arg Leu Asn Ser Thr His Leu Tyr 25 Ala Cys Gly Thr His Ala Phe Gln Pro Leu Cys Ala Ala Ile Asp Ala 40 Glu Ala Phe Thr Leu Pro Thr Ser Phe Glu Glu Gly Lys Glu Lys Cys 55 60 Pro Tyr Asp Pro Ala Arg Gly Phe Thr Gly Leu Ile Ile Asp Gly Gly 70 75 Leu Tyr Thr Ala Thr Arg Tyr Glu Phe Arg Ser Ile Pro Asp Ile Arg 85 90 Arg Ser Arg His Pro His Ser Leu Arg Thr Glu Glu Thr Pro Met His 100 105 110 Trp Leu Asn Asp Ala Glu Phe Val Phe Ser Val Leu Val Arg Glu Ser 115 120 125 Lys Ala Ser Ala Val Gly Asp Asp Lys Val Tyr Tyr Phe Phe Thr 135 Glu Arg Ala Thr Glu Glu Gly Ser Gly Ser Phe Thr Gln Ser Arg Ser 150 160 155 Ser His Arg Val Ala Arg Val Ala Arg Val Cys Lys Gly Asp Leu Gly 170 Gly Lys Lys Ile Leu Gln Lys Lys Trp Thr Ser Phe Leu Lys Ala Arg

			180					185					190		
Leu	Ile	Cys 195	His	Ile	Pro	Leu	Tyr 200	Glu	Thr	Leu	Arg	Gly 205	Val	Cys	Ser
	Asp 210					215					220				
225	Ser				230					235					240
	Leu			245					250					255	
	Asp		260					265					270		
	Arg	275					280					285			
	Ser 290					295					300				
305	Pro				310					315					320
	Leu			325					330					335	
	Thr		340					345					350		
	Gly	355					360					365			
	Glu 370					375					380				
385	Ile				390					395					400
	Ile			405					410					415	-
	Cys		420					425	_	_	_	_	430	_	
	Ala	435					440					445			
	Thr 450					455					460				
465	Ser				470					475					480
	Pro			485					490					495	
	Val		500					505					510		
	Glu Leu	515					520					525			
	530 Gln					535					540				
545	Leu				550					555					560
				565					570					575	
	His Pro		580					585			AId	ьeu	590	ASII	GTII
110	110	595	DGT	110	11P	SET	600	ner	ALG	vaı					

<210> 40 <211> 1797

```
<212> DNA
<213> homo sapiens
<400> 40
atgcaaagca aatgtcatca aaaagggaaa aacaaccaga cggagtgctt taaccatgtg
                                                                        60
cggttcctgc agcggctcaa ttctacccac ctctatgcat gtgggactca cgccttccag
                                                                       120
cccctctgtg cagccattga tgctgaggcc ttcaccttgc caaccagctt cgaggaggg
                                                                       180
aaggagaagt gtccttatga cccaqccqt qqcttcacaq qcctcatcat tqatqqaqqc
                                                                       240
ctctacacag ccactaggta tgaattccgg agcattcctg acatccgccg gagccgccac
                                                                       300
ccacactccc tgagaactga ggagacacca atgcattggc tcaatgatgc ggagtttgtg
                                                                       360
ttctccgtcc tcgtgcggga gagcaaggcc agtgcagtgg gtgatgatga caaggtgtac
                                                                       420
tacttettea eggagegtge caetgaggag ggetetggea getteactea gageegeage
                                                                       480
agtcaccgtg tggcccgtgt ggctcgygtc tgcaagggag acctgggagg gaagaagatc
                                                                       540
ctgcagaaga agtggacttc cttcctgaaa gcccgtctca tctgccacat tccactgtat
                                                                       600
gagacactgc gtggggtctg cagcctggat gctgaaacct caagccgtac acacttctat
                                                                       660
gcagccttca cgctgagcac acagtggaag accctggagg cctcagccat ctgccgctat
                                                                       720
gacctggcag agatccaggc tgtctttgca ggaccctata tggaatacca ggatggttcc
                                                                       780
cggcgctggg gtcgctatga gggtggggtg cctgagcccc ggcctggctc gtgtatcaca
                                                                       840
gattcattgc gcagccaagg ctacaattca tcccaagact tgccatccct ggtcctggac
                                                                       900
tttgtaaagt tgcacccact gatggctcgg cccgttgtgc ccacacgtgg acggccctg
                                                                       960
ctgctcaagc gcaacatacg ctacacacac cttacaggga cacctgtcac cacgcctgct
                                                                      1020
ggacctacct atgacctgct ctttctgggc acagctgatg gctggatcca caaggccgta
                                                                      1080
gtcctgggct ctgggatgca cattattgaa gagacacaag tgttcaggga gtcccagtct
                                                                      1140
gtggaaaatc tagtcatctc tctattgcag cacagcctct atgtgggggc tcctagcgga
                                                                      1200
gtcatccage taccactete cagetgetee egetacegat cetgetatga etgeatettg
                                                                      1260
gcccgagacc cctactgtgg ctgggaccct ggcacccatg cctgcgcagc agccaccacc
                                                                      1320
atagccaaca ggacagcact gatacaggac atagagagag gaaatcgagg ctgtgagagc
                                                                      1380
agcagggata caggcagggc tctgcaggtc catatgggct caatgtcacc accctctgca
                                                                      1440
tggccctgtg tgctggatgg tcctgaaacc agacaagtcc tctgccagcc acctaagccc
                                                                      1500
tgcgtacatt cacatgcaca catggaagaa tgtttatcgg ctgggctgca gtgccccac
                                                                      1560
cctcaccttc tcctggtgca ttcttgtttc atccctgctt ctggacttgg ggtaccctcc
                                                                      1620
caattgccac atcctatctg gtcctcttcc ccagccccat gtggtgacct ctttgtcaag
                                                                      1680
agettgggaa egggeeagee tggggaggta agaetgeate acteeectee teteeettee
                                                                      1740
tgtgtggccc ttgtgaatca gcctccccac tctccttggt cattctcaag agtatga
                                                                      1797
<210> 41
<211> 598
<212> PRT
<213> homo sapiens
<400> 41
Met Gln Ser Lys Cys His Gln Lys Gly Lys Asn Asn Gln Thr Glu Cys
1
                                    10
Phe Asn His Val Arg Phe Leu Gln Arg Leu Asn Ser Thr His Leu Tyr
                                25
Ala Cys Gly Thr His Ala Phe Gln Pro Leu Cys Ala Ala Ile Asp Ala
                            40
Glu Ala Phe Thr Leu Pro Thr Ser Phe Glu Glu Gly Lys Glu Lys Cys
                        55
Pro Tyr Asp Pro Ala Arg Gly Phe Thr Gly Leu Ile Ile Asp Gly Gly
                    70
                                        75
```

90

Leu Tyr Thr Ala Thr Arg Tyr Glu Phe Arg Ser Ile Pro Asp Ile Arg

Arg Ser Arg His Pro His Ser Leu Arg Thr Glu Glu Thr Pro Met His

105 Trp Leu Asn Asp Ala Glu Phe Val Phe Ser Val Leu Val Arg Glu Ser

		115					120					125			
Lys	Ala 130	Ser	Ala	Val	Gly	Asp 135	Asp	Asp	Lys	Val	Tyr 140		Phe	Phe	Thr
Glu 145	Arg	Ala	Thr	Glu	Glu 150	Gly	Ser	Gly	Ser	Phe 155	Thr	G1n	Ser	Arg	Ser 160
Ser	His	Arg	Val	Ala 165	Arg	Val	Ala	Arg	Val 170	Cys	Lys	Gly	Asp	Leu 175	Gly
Gly	Lys	Lys	Ile 180	Leu	Gln	Lys	Lys	Trp 185	Thr	Ser	Phe	Leu	Lys 190	Ala	Arg
Leu	Ile	Суs 195	His	Ile	Pro	Leu	Tyr 200	Glu	Thr	Leu	Arg	Gly 205	Val	Cys	Ser
Leu	Asp 210	Ala	Glu	Thr	Ser	Ser 215	Arg	Thr	His	Phe	Tyr 220	Ala	Ala	Phe	Thr
Leu 225	Ser	Thr	Gln	Trp	Lys 230	Thr	Leu	Glu	Ala	Ser 235	Ala	Ile	Cys	Arg	Tyr 240
Asp	Leu	Ala	Glu	Ile 245	Gln	Ala	Va1	Phe	Ala 250	Gly	Pro	Tyr	Met	G1u 255	Tyr
			260		Arg			265					270		
		275			Суз		280					285			
	290				Leu	295					300			_	
305					Arg 310					315	_				320
				325	Ile				330					335	
			340		Pro			345					350		
Asp	Gly	Trp 355	Ile	His	Lys	Ala	Val 360	Val	Leu	Gly	Ser	Gly 365	Met	His	Ile
	370				Val	375	-				380				
385					Gln 390					395					400
				405	Leu				410					415	
			420		Arg			425					430		
		435			Ala		440					445			
	450				Gly	455					460				
465					Val 470					475					480
				485	Asp				490					495	
			500		Val			505					510	_	
		515			Cys		520					525			
	530				Ser	535					540				
545					Ser 550					555					560
ser	ьeu	GТĀ	ınr	GТĀ	Gln	Pro	GTÀ	Glu	val	Arg	Leu	His	His	Ser	Pro

```
565
                                     570
                                                         575
Pro Leu Pro Ser Cys Val Ala Leu Val Asn Gln Pro Pro His Ser Pro
                                585
Trp Ser Phe Ser Arg Val
        595
<210> 42
<211> 2235
<212> DNA
<213> homo sapiens
<400> 42
atgcaaagca aatgtcatca aaaagggaaa aacaaccaga cggagtgctt taaccatgtg
                                                                        60
eggtteetge ageggeteaa ttetacceae etetatgeat gtgggaetea egeetteeag
                                                                       120
cccctctgtg cagccattga tgctgaggcc ttcaccttgc caaccagctt cgaggagggg
                                                                       180
aaggagaagt gtccttatga cccagcccgt ggcttcacag gcctcatcat tgatggaggc
                                                                       240
ctctacacag ccactaggta tgaattccgg agcattcctg acatccgccg gagccgccac
                                                                       300
ccacactccc tgagaactga ggagacacca atgcattggc tcaatgatgc ggagtttgtg
                                                                       360
ttctccgtcc tcgtgcggga gagcaaggcc agtgcagtgg gtgatgatga caaggtgtac
                                                                       420
tacttettea eggagegtge eactgaggag ggetetggea getteaetea gageegeage
                                                                       480
agtcaccgtg tggcccgtgt ggctcgygtc tgcaagggag acctgggagg gaagaagatc
                                                                       540
ctgcagaaga agtggacttc cttcctgaaa gcccgtctca tctgccacat tccactgtat
                                                                       600
gagacactgc gtggggtctg cagcctggat gctgaaacct caagccgtac acacttctat
                                                                       660
gcagcettea egetgageae acagtggaag accetggagg ceteageeat etgeegetat
                                                                       720
gacctggcag agatccaggc tgtctttgca ggaccctata tggaatacca ggatggttcc
                                                                       780
cggcgctggg gtcgctatga gggtggggtg cctgagcccc ggcctggctc gtgtatcaca
                                                                       840
gattcattgc gcagccaagg ctacaattca tcccaagact tgccatccct ggtcctggac
                                                                       900
                                                                       960
tttgtaaagt tgcacccact gatggctcgg cccgttgtgc ccacacgtgg acggcccctg
ctgctcaagc gcaacatacg ctacacacac cttacaggga cacctgtcac cacqcctgct
                                                                      1020
ggacctacct atgacctgct ctttctgggc acagctgatg gctggatcca caaggccgta
                                                                      1080
gtcctgggct ctgggatgca cattattgaa gagacacaag tgttcaggga gtcccagtct
                                                                      1140
gtggaaaatc tagtcatctc tctattgcag cacagcctct atgtgggggc tcctagcgga
                                                                      1200
gtcatccagc taccactctc cagetgetee egetacegat ectgetatga etgeatettg
                                                                      1260
gcccgagacc cctactgtgg ctgggaccct ggcacccatg cctgcgcagc agccaccacc
                                                                      1320
atagccaaca ggtcccaggg aagcaggaca gcactgatac aggacataga gagaggaaat
                                                                      1380
cgaggctgtg agagcagcag ggatacaggg ccaccaccac cactgaagac ccgctctgtg
                                                                      1440
ctccggggtg atgatgtcct cctgccctgt gaccagccat ccaacctggc ccgggccttg
                                                                      1500
tggctactca atgggagcat gggcctgagc gatgggcagg gtggctaccg tgtgggcgtg
                                                                      1560
gacgggctgc tggttacaga tgcacagcct gagcacagtg gcaactatgg ctgctatgcc
                                                                      1620
gaggaaaatg gcctccgcac cctgctggcc tcctatagtc tcacagtccg gccagccact
                                                                      1680
cctgccccag ctccaaaagc ccctgccaca cctggggcac agctggcacc tgatgtgaga
                                                                      1740
ctgctctatg tgctagccat tgccgcgctt ggtggccyct gcctcatcct ggcctcctcc
                                                                      1800
ctcctctatg tggcctgtct gcgggaaggc agacgagggc gccgacggaa atactcactg
                                                                     1860
ggtcgggcca gccgggcagg aggatctgcg gtgcaactgc agacagtctc aggccagtgt
                                                                      1920
cctggagagg aagatgaggg tgatgatgag ggggctgggg gcctggaggg cagctgtctc
                                                                     1980
cagatcatcc ctggggaggg agccccagcc ccaccaccc caccgccccc accgccaccg
                                                                      2040
gctgagctga ccaatggctt ggtggcactg cccagccggc tgcggaggat gaatgqcaat
                                                                      2100
agctatgtgc ttctgaggca gagcaacaat ggagtaccag cagggccctg ctccttcgcc
                                                                      2160
gaggaactca gccgcatcct ggaaaaaagg aagcacacgc agctcgtgga gcagctagat
                                                                      2220
gagagetetg tetga
                                                                      2235
<210> 43
<211> 744
<212> PRT
<213> homo sapiens
```

<400> 43 Met Gln Ser Lys Cys His Gln Lys Gly Lys Asn Asn Gln Thr Glu Cys 10 Phe Asn His Val Arg Phe Leu Gln Arg Leu Asn Ser Thr His Leu Tyr Ala Cys Gly Thr His Ala Phe Gln Pro Leu Cys Ala Ala Ile Asp Ala 40 Glu Ala Phe Thr Leu Pro Thr Ser Phe Glu Glu Gly Lys Glu Lys Cys 55 Pro Tyr Asp Pro Ala Arg Gly Phe Thr Gly Leu Ile Ile Asp Gly Gly 70 75 Leu Tyr Thr Ala Thr Arg Tyr Glu Phe Arg Ser Ile Pro Asp Ile Arg 90 Arg Ser Arg His Pro His Ser Leu Arg Thr Glu Glu Thr Pro Met His 100 105 Trp Leu Asn Asp Ala Glu Phe Val Phe Ser Val Leu Val Arg Glu Ser 120 Lys Ala Ser Ala Val Gly Asp Asp Asp Lys Val Tyr Tyr Phe Phe Thr 135 140 Glu Arg Ala Thr Glu Glu Gly Ser Gly Ser Phe Thr Gln Ser Arg Ser 155 150 Ser His Arg Val Ala Arg Val Ala Arg Val Cys Lys Gly Asp Leu Gly 170 Gly Lys Lys Ile Leu Gln Lys Lys Trp Thr Ser Phe Leu Lys Ala Arg 185 180 Leu Ile Cys His Ile Pro Leu Tyr Glu Thr Leu Arg Gly Val Cys Ser 200 205 Leu Asp Ala Glu Thr Ser Ser Arg Thr His Phe Tyr Ala Ala Phe Thr 215 220 Leu Ser Thr Gln Trp Lys Thr Leu Glu Ala Ser Ala Ile Cys Arg Tyr 230 235 Asp Leu Ala Glu Ile Gln Ala Val Phe Ala Gly Pro Tyr Met Glu Tyr 245 250 Gln Asp Gly Ser Arg Arg Trp Gly Arg Tyr Glu Gly Gly Val Pro Glu 260 265 270 Pro Arg Pro Gly Ser Cys Ile Thr Asp Ser Leu Arg Ser Gln Gly Tyr 280 285 Asn Ser Ser Gln Asp Leu Pro Ser Leu Val Leu Asp Phe Val Lys Leu 300 295 His Pro Leu Met Ala Arg Pro Val Val Pro Thr Arg Gly Arg Pro Leu 315 310 Leu Leu Lys Arg Asn Ile Arg Tyr Thr His Leu Thr Gly Thr Pro Val 325 330 Thr Thr Pro Ala Gly Pro Thr Tyr Asp Leu Leu Phe Leu Gly Thr Ala 345 Asp Gly Trp Ile His Lys Ala Val Val Leu Gly Ser Gly Met His Ile 360 365 355 Ile Glu Glu Thr Gln Val Phe Arg Glu Ser Gln Ser Val Glu Asn Leu 375 Val Ile Ser Leu Leu Gln His Ser Leu Tyr Val Gly Ala Pro Ser Gly 395 390 Val Ile Gln Leu Pro Leu Ser Ser Cys Ser Arg Tyr Arg Ser Cys Tyr 410 Asp Cys Ile Leu Ala Arg Asp Pro Tyr Cys Gly Trp Asp Pro Gly Thr 425 His Ala Cys Ala Ala Ala Thr Thr Ile Ala Asn Arg Ser Gln Gly Ser

```
435
                        440
                                          445
Arg Thr Ala Leu Ile Gln Asp Ile Glu Arg Gly Asn Arg Gly Cys Glu
                    455
Ser Ser Arg Asp Thr Gly Pro Pro Pro Pro Leu Lys Thr Arg Ser Val
       470
                                  475
Leu Arg Gly Asp Asp Val Leu Leu Pro Cys Asp Gln Pro Ser Asn Leu
             485
                   490
Ala Arg Ala Leu Trp Leu Leu Asn Gly Ser Met Gly Leu Ser Asp Gly
                            505
Gln Gly Gly Tyr Arg Val Gly Val Asp Gly Leu Leu Val Thr Asp Ala
                         520
Gln Pro Glu His Ser Gly Asn Tyr Gly Cys Tyr Ala Glu Glu Asn Gly
                     535
Leu Arg Thr Leu Leu Ala Ser Tyr Ser Leu Thr Val Arg Pro Ala Thr
                 550
                                   555
Pro Ala Pro Ala Pro Lys Ala Pro Ala Thr Pro Gly Ala Gln Leu Ala
             565
                               570
Pro Asp Val Arg Leu Leu Tyr Val Leu Ala Ile Ala Ala Leu Gly Gly
         580 585
Leu Cys Leu Ile Leu Ala Ser Ser Leu Leu Tyr Val Ala Cys Leu Arg
            600 605
Glu Gly Arg Arg Gly Arg Arg Lys Tyr Ser Leu Gly Arg Ala Ser
                    615
Arg Ala Gly Gly Ser Ala Val Gln Leu Gln Thr Val Ser Gly Gln Cys
                 630
                                   635
Pro Gly Glu Glu Asp Glu Gly Asp Asp Glu Gly Ala Gly Gly Leu Glu
                                650
Gly Ser Cys Leu Gln Ile Ile Pro Gly Glu Gly Ala Pro Ala Pro Pro
   660
                           665
Pro Pro Pro Pro Pro Pro Pro Ala Glu Leu Thr Asn Gly Leu Val
                        680
Ala Leu Pro Ser Arg Leu Arg Arg Met Asn Gly Asn Ser Tyr Val Leu
                                       700
       695
Leu Arg Gln Ser Asn Asn Gly Val Pro Ala Gly Pro Cys Ser Phe Ala
     710
                                  715
Glu Glu Leu Ser Arg Ile Leu Glu Lys Arg Lys His Thr Gln Leu Val
             725
                             730
Glu Gln Leu Asp Glu Ser Ser Val
          740
```

<211> 2220

<212> DNA

<213> homo sapiens

## <400> 44

atgcaaagca	aatgtcatca	aaaagggaaa	aacaaccaga	cggagtgctt	taaccatgtg	60
cggttcctgc	agcggctcaa	ttctacccac	ctctatgcat	gtgggactca	cgccttccag	120
cccctctgtg	cagccattga	tgctgaggcc	ttcaccttgc	caaccagctt	cgaggagggg	180
aaggagaagt	gtccttatga	cccagcccgt	ggcttcacag	gcctcatcat	tgatggaggc	240
ctctacacag	ccactaggta	tgaattccgg	agcattcctg	acatccgccg	gagccgccac	300
ccacactccc	tgagaactga	ggagacacca	atgcattggc	tcaatgatgc	ggagtttgtg	360
ttctccgtcc	tcgtgcggga	gagcaaggcc	agtgcagtgg	gtgatgatga	caaggtgtac	420
tacttcttca	cggagcgtgc	cactgaggag	ggctctggca	gcttcactca	gagccgcagc	480
agtcaccgtg	tggcccgtgt	ggctcgygtc	tgcaagggag	acctgggagg	gaagaagatc	540
ctgcagaaga	agtggacttc	cttcctgaaa	gcccgtctca	tctgccacat	tccactgtat	600

```
Ala Val Gln Leu Gln Thr Val Ser Gly Gln Cys Pro Gly Glu Glu Asp
                    630
                                        635
Glu Gly Asp Asp Glu Gly Ala Gly Gly Leu Glu Gly Ser Cys Leu Gln
                                    650
                645
Ile Ile Pro Gly Glu Gly Ala Pro Ala Pro Pro Pro Pro Pro Pro
                                665
Pro Pro Pro Ala Glu Leu Thr Asn Gly Leu Val Ala Leu Pro Ser Arg
                                                 685
                            680
Leu Arg Arg Met Asn Gly Asn Ser Tyr Val Leu Leu Arg Gln Ser Asn
                        695
                                             700
Asn Gly Val Pro Ala Gly Pro Cys Ser Phe Ala Glu Glu Leu Ser Arg
                                                             720
                    710
                                        715
Ile Leu Glu Lys Arg Lys His Thr Gln Leu Val Glu Gln Leu Asp Glu
                                    730
                725
```

Ser Ser Val

<210> 46 <211> 2316 <212> DNA

<213> homo sapiens

<400> 46

```
60
atgcaaagca aatgtcatca aaaagggaaa aacaaccaga cggagtgctt taaccatgtg
cggttcctgc agcggctcaa ttctacccac ctctatgcat gtgggactca cgccttccag
                                                                       120
                                                                       180
cccctctgtg cagccattga tgctgaggcc ttcaccttgc caaccagctt cgaggagggg
                                                                       240
aaggagaagt gtccttatga cccagcccgt ggcttcacag gcctcatcat tgatggaggc
                                                                       300
ctctacacag ccactaggta tgaattccgg agcattcctg acatccgccg gagccgccac
ccacactccc tgagaactga ggagacacca atgcattggc tcaatgatgc ggagtttgtg
                                                                       360
                                                                       420
ttctccgtcc tcgtgcggga gagcaaggcc agtgcagtgg gtgatgatga caaggtgtac
                                                                       480
tacttcttca cggagcgtgc cactgaggag ggctctggca gcttcactca gagccgcagc
agtcaccgtg tggcccgtgt ggctcgygtc tgcaagggag acctgggagg gaagaagatc
                                                                       540
ctgcagaaga agtggacttc cttcctgaaa gcccgtctca tctgccacat tccactgtat
                                                                       600
                                                                       660
qaqacactqc qtqqqqtctq cagcctggat gctgaaacct caagccgtac acacttctat
gcagccttca cgctgagcac acagtggaag accctggagg cctcagccat ctgccgctat
                                                                       720
gacctggcag agatccaggc tgtctttgca ggaccctata tggaatacca ggatggttcc
                                                                       780
cggcgctggg gtcgctatga gggtggggtg cctgagcccc ggcctggctc gtgtatcaca
                                                                       840
gattcattgc gcagccaagg ctacaattca tcccaagact tgccatccct ggtcctggac
                                                                       900
tttgtaaagt tgcacccact gatggctcgg cccgttgtgc ccacacgtgg acggcccctg
                                                                       960
ctgctcaagc gcaacatacg ctacacacac cttacaggga cacctgtcac cacgcctgct
                                                                      1020
                                                                      1080
ggacctacct atgacctgct ctttctgggc acagctgatg gctggatcca caaggccgta
gtcctgggct ctgggatgca cattattgaa gagacacaag tgttcaggga gtcccagtct
                                                                      1140
                                                                      1200
gtggaaaatc tagtcatctc tctattgcag cacagcctct atgtgggggc tcctagcgga
gtcatccagc taccactctc cagctgctcc cgctaccgat cctgctatga ctgcatcttg
                                                                      1260
                                                                      1320
qcccqaqacc cctactgtgg ctgggaccct ggcacccatg cctgcgcagc agccaccacc
                                                                      1380
atagccaaca ggtcccaggg aagcaggaca gcactgatac aggacataga gagaggaaat
                                                                      1440
cqaggctqtg agagcagcag ggatacaggg ccaccaccac cactgaagac ccgctctgtg
                                                                      1500
ctccggggtg atgatgtcct cctgccctgt gaccagccat ccaacctggc ccgggccttg
tggctactca atgggagcat gggcctgagc gatgggcagg gtggctaccg tgtgggcgtg
                                                                      1560
                                                                      1620
gacgggctgc tggttacaga tgcacagcct gagcacagtg gcaactatgg ctgctatgcc
gaggaaaatg gcctccgcac cctgctggcc tcctatagtc tcacagtccg gccagccact
                                                                      1680
                                                                      1740
cctqccccaq ctccaaaaagc ccctqccaca cctggggcac agctggcacc tgatgtgaga
                                                                      1800
ctgctctatq tgctagccat tgccgcgctt ggtggccyct gcctcatcct ggcctcctcc
                                                                      1860
ctcctctatg tggcctgtct gcgggaaggc agacgagggc gccgacggaa atactcactg
                                                                      1920
ggtcgggcca gccgggcagg aggatctgcg gtgcaactgc agacagtctc aggcagggct
ctgcaggtcc atatgggctc aatgtcacca ccctctgcat ggccctgtgt gctggatggt
                                                                      1980
```

cctgaaacca gacaagteet etgecageca cctaageeet gegtacatte acatgeacac atggaagaat gtttategge tgggetgeag tgececeace eteacettet eetggtgeat tettgttea teeetgette tggaettggg gtaceeteee aattgecaca teetatetgg teeteetteee cageeeeatg tggtgacete tttgteaaga gettgggaac gggecageet ggggaggtaa gactgeatea eteeeteet eteeetteet gtgtggeeet tgtgaateag eeteeeede eteettggte atteteaaga gtatga	2040 2100 2160 2220 2280 2316
<210> 47 <211> 771 <212> PRT <213> homo sapiens	
<400> 47	
Met Gln Ser Lys Cys His Gln Lys Gly Lys Asn Asn Gln Thr Glu Cys  1 10 15	
Phe Asn His Val Arg Phe Leu Gln Arg Leu Asn Ser Thr His Leu Tyr 20 25 30	
Ala Cys Gly Thr His Ala Phe Gln Pro Leu Cys Ala Ala Ile Asp Ala	

Glu Ala Phe Thr Leu Pro Thr Ser Phe Glu Glu Gly Lys Glu Lys Cys Pro Tyr Asp Pro Ala Arg Gly Phe Thr Gly Leu Ile Ile Asp Gly Gly Leu Tyr Thr Ala Thr Arg Tyr Glu Phe Arg Ser Ile Pro Asp Ile Arg Arg Ser Arg His Pro His Ser Leu Arg Thr Glu Glu Thr Pro Met His Trp Leu Asn Asp Ala Glu Phe Val Phe Ser Val Leu Val Arg Glu Ser Lys Ala Ser Ala Val Gly Asp Asp Asp Lys Val Tyr Tyr Phe Phe Thr Glu Arg Ala Thr Glu Glu Gly Ser Gly Ser Phe Thr Gln Ser Arg Ser Ser His Arg Val Ala Arg Val Ala Arg Val Cys Lys Gly Asp Leu Gly Gly Lys Lys Ile Leu Gln Lys Lys Trp Thr Ser Phe Leu Lys Ala Arg Leu Ile Cys His Ile Pro Leu Tyr Glu Thr Leu Arg Gly Val Cys Ser Leu Asp Ala Glu Thr Ser Ser Arg Thr His Phe Tyr Ala Ala Phe Thr Leu Ser Thr Gln Trp Lys Thr Leu Glu Ala Ser Ala Ile Cys Arg Tyr Asp Leu Ala Glu Ile Gln Ala Val Phe Ala Gly Pro Tyr Met Glu Tyr Gln Asp Gly Ser Arg Arg Trp Gly Arg Tyr Glu Gly Gly Val Pro Glu Pro Arg Pro Gly Ser Cys Ile Thr Asp Ser Leu Arg Ser Gln Gly Tyr Asn Ser Ser Gln Asp Leu Pro Ser Leu Val Leu Asp Phe Val Lys Leu His Pro Leu Met Ala Arg Pro Val Val Pro Thr Arg Gly Arg Pro Leu Leu Leu Lys Arg Asn Ile Arg Tyr Thr His Leu Thr Gly Thr Pro Val Thr Thr Pro Ala Gly Pro Thr Tyr Asp Leu Leu Phe Leu Gly Thr Ala

			340					345					350		
Agn	Gly	Trn		Hic	Tare	Δl =	T.C. 7		Ten	C132	Cor	Clar		Wic	Tla
nap	GTĀ	355	116	111.5	гуу	пта	360	vai	теи	GTA	ser	365	Mec	птъ	TTE
Tle	Glu		Thr	Gln	Val	Phe		Glu	Ser	Gln	Ser		Glu	Asn	Len
110	370	O.L. G.		0111	, 41	375	9	J_ u	201	0111	380	V CL	O.L.	11011	200
Val	Ile	Ser	Leu	Leu	Gln		Ser	Leu	Tvr	Val		Ala	Pro	Ser	Glv
385					390				- 4 -	395	2				400
Val	Ile	G1n	Leu	Pro	Leu	Ser	Ser	Cys	Ser	Arg	Tyr	Arg	Ser	Cys	Tyr
				405					410					415	
Asp	Суѕ	Ile	Leu 420	Ala	Arg	Asp	Pro	Tyr 425	Cys	Gly	Trp	Asp	Pro 430	Gly	Thr
His	Ala	Cys 435	Ala	Ala	Ala	Thr	Thr 440		Ala	Asn	Arg	Ser		Gly	Ser
Δτα	Thr		T.611	T1_	Gln	Δen		Glu	Δrα	Clv	Acn		Clv	Cve	G111
	450					455			_		460	-	_	_	
	Ser	Arg	Asp	Thr		Pro	Pro	Pro	Pro		Lys	Thr	Arg	Ser	
465	7.20	0111	7 an	7 ~~	470	T 011	T 011	Dres	Crea	475	~1~	Drag	(10.00	70	480
ьeu	Arg	GIÀ	Asp	485	Val	ьеu	Leu	PIO	490	Asp	GIII	Pro	ser	495	Leu
Ala	Arg	Ala	Leu 500	Trp	Leu	Leu	Asn	Gly 505	Ser	Met	Gly	Leu	Ser 510	Asp	Gly
Gln	Gly	Gly 515	Tyr	Arg	Val	Gly	Val 520	Asp	Gly	Leu	Leu	Val 525	Thr	Asp	Ala
G1n	Pro	Glu	His	Ser	Gly	Asn 535		Gly	Cys	Tyr	Ala 540		Glu	Asn	Gly
T <sub>(</sub> e)1	Arg	Thr	T.e11	T.e.11	Δla		Tur	Ser	T.em	Thr		Ara	Pro	Δla	Thr
545					550					555					560
Pro	Ala	Pro	Ala	Pro 565	Lys	Ala	Pro	Ala	Thr 570	Pro	GIĀ	Ala	Gln	Leu 575	Ala
Pro	Asp	Val	Arg 580	Leu	Leu	Tyr	Val	Leu 585	Ala	Ile	Ala	Ala	Leu 590	Gly	Gly
Leu	Cys	Leu 595	Ile	Leu	Ala	Ser	Ser 600	Leu	Leu	Tyr	Val	Ala 605	Cys	Leu	Arg
Glu	Gly 610	Arg	Arg	Gly	Arg	Arg 615	Arg	Lys	Tyr	Ser	Leu 620	Gly	Arg	Ala	Ser
	Ala	Gly	Gly	Ser			Gln	Leu	Gln			Ser	Gly	Arg	Ala
625	07	777	TT.	3.6 - I-	630	<b>G</b> -	36 1	<b>a</b> .	ъ	635	~		-	_	640
neu	Gln	Val	nis	Met 645	GTĀ	ser	мес	ser	650	Pro	ser	Ala	Trp	655	Суѕ
Val	Leu	Asp	Gly 660	Pro	Glu	Thr	Arg	Gln 665	Val	Leu	Cys	Gln	Pro 670	Pro	Lys
Pro	Cys	Val 675	His	Ser	His	Ala	His 680	Met	Glu	Glu	Cys	Leu 685	Ser	Ala	Gly
Leu	Gln 690		Pro	His	Pro	His		Leu	Leu	Val	His		Cys	Phe	Ile
Pro	Ala	Ser	Glv	T. 🗕 11	G1v		Pro	Car	Gln	T 011		Шi с	Pro	T10	Trn
705					710					715					720
Ser	Ser	Ser	Pro	Ala 725	Pro	Cys	Gly	Asp	Leu 730	Phe	Val	Lys	Ser	Leu 735	Gly
Thr	Gly	Gln	Pro 740	Gly	Glu	Val	Arg	Leu 745	His	His	Ser	Pro	Pro 750	Leu	Pro
Ser	Cys	Val 755		Leu	Val	Asn	Gln 760		Pro	His	Ser	Pro 765		Ser	Phe
Ser	Arg						, 50					, , , ,			
	770														

```
<211> 2301
<212> DNA
<213> homo sapiens
<400> 48
atgcaaagca aatgtcatca aaaagggaaa aacaaccaga cggagtgctt taaccatgtg
                                                                        60
                                                                       120
cggttcctgc agcggctcaa ttctacccac ctctatgcat gtgggactca cgccttccag
cccctctgtg cagccattga tgctgaggcc ttcaccttgc caaccagctt cgaggagggg
                                                                       180
                                                                       240
aaggagaagt gteettatga eecageeegt ggetteacag geeteateat tgatggagge
                                                                       300
ctctacacag ccactaggta tgaattccgg agcattcctg acatccgccg gagccgccac
                                                                       360
ccacactccc tgagaactga ggagacacca atgcattggc tcaatgatgc ggagtttgtg
                                                                       420
ttctccgtcc tcgtgcggga gagcaaggcc agtgcagtgg gtgatgatga caaggtgtac
                                                                       480
tacttettea eggagegtge eactgaggag ggetetggea getteactea gageegeage
                                                                       540
agtcaccgtg tggcccgtgt ggctcgygtc tgcaagggag acctgggagg gaagaagatc
ctgcagaaga agtggacttc cttcctgaaa gcccgtctca tctgccacat tccactgtat
                                                                       600
gagacactgc gtggggtctg cagcctggat gctgaaacct caagccgtac acacttctat
                                                                       660
                                                                       720
gcagccttca cgctgagcac acagtggaag accctggagg cctcagccat ctgccgctat
                                                                       780
gacctggcag agatccaggc tgtctttgca ggaccctata tggaatacca ggatggttcc
                                                                       840
cggcgctggg gtcgctatga gggtggggtg cctgagcccc ggcctggctc gtgtatcaca
                                                                       900
gattcattgc gcagccaagg ctacaattca tcccaagact tgccatccct ggtcctggac
                                                                       960
tttgtaaagt tgcacccact gatggctcgg cccgttgtgc ccacacgtgg acggccctg
                                                                      1020
ctgctcaagc gcaacatacg ctacacacac cttacaggga cacctgtcac cacgcctgct
                                                                      1080
ggacctacct atgacctgct ctttctgggc acagctgatg gctggatcca caaggccgta
                                                                      1140
gtcctgggct ctgggatgca cattattgaa gagacacaag tgttcaggga gtcccagtct
gtggaaaatc tagtcatctc tctattgcag cacagcctct atgtgggggc tcctagcgga
                                                                      1200
gtcatccagc taccactctc cagctgctcc cgctaccgat cctgctatga ctgcatcttg
                                                                      1260
                                                                      1320
gcccgagacc cctactgtgg ctgggaccct ggcacccatg cctgcgcagc agccaccacc
                                                                      1380
atagccaaca ggacagcact gatacaggac atagagagag gaaatcgagg ctgtgagagc
                                                                      1440
agcagggata cagggccacc accaccactg aagacccgct ctgtgctccg gggtgatgat
                                                                      1500
gtcctcctgc cctgtgacca gccatccaac ctggcccggg ccttgtggct actcaatggg
agcatgggcc tgagcgatgg gcagggtggc taccgtgtgg gcgtggacgg gctgctggtt
                                                                      1560
acagatgcac agcctgagca cagtggcaac tatggctgct atgccgagga aaatggcctc
                                                                      1620
cgcaccetge tggcctceta tagtetcaca gtccggccag ccactcetge cccagetcca
                                                                      1680
aaagcccctg ccacacctgg ggcacagctg gcacctgatg tgagactgct ctatgtgcta
                                                                      1740
                                                                      1800
gccattgccg cgcttggtgg ccyctgcctc atcctggcct cctcctcct ctatgtggcc
                                                                      1860
tgtctgcggg aaggcagacg agggcgccga cggaaatact cactgggtcg ggccagccgg
                                                                      1920
gcaggaggat ctgcggtgca actgcagaca gtctcaggca gggctctgca ggtccatatg
ggctcaatgt caccacctc tgcatggccc tgtgtgctgg atggtcctga aaccagacaa
                                                                      1980
gtcctctgcc agccacctaa gccctgcgta cattcacatg cacacatgga agaatgttta
                                                                      2040
teggetggge tgeagtgeec ceaeceteac etteteetgg tgeattettg ttteateeet
                                                                      2100
                                                                      2160
gcttctggac ttggggtacc ctcccaattg ccacatccta tctggtcctc ttccccagcc
                                                                      2220
ccatgtggtg acctetttgt caagagettg ggaacgggec ageetgggga ggtaagaetg
                                                                      2280
catcactece etecteteec tteetgtgtg geeettgtga ateageetee ecaeteteet
tggtcattct caagagtatg a
                                                                      2301
<210> 49
<211> 766
<212> PRT
<213> homo sapiens
<400> 49
Met Gln Ser Lys Cys His Gln Lys Gly Lys Asn Asn Gln Thr Glu Cys
                                    10
Phe Asn His Val Arg Phe Leu Gln Arg Leu Asn Ser Thr His Leu Tyr
            20
                                2.5
                                                    30
```

Ala Cys Gly Thr His Ala Phe Gln Pro Leu Cys Ala Ala Ile Asp Ala 40 Glu Ala Phe Thr Leu Pro Thr Ser Phe Glu Glu Gly Lys Glu Lys Cys 55 Pro Tyr Asp Pro Ala Arg Gly Phe Thr Gly Leu Ile Ile Asp Gly Gly 75 Leu Tyr Thr Ala Thr Arg Tyr Glu Phe Arg Ser Ile Pro Asp Ile Arg 90 85 Arg Ser Arg His Pro His Ser Leu Arg Thr Glu Glu Thr Pro Met His 105 100 Trp Leu Asn Asp Ala Glu Phe Val Phe Ser Val Leu Val Arg Glu Ser 120 125 Lys Ala Ser Ala Val Gly Asp Asp Asp Lys Val Tyr Tyr Phe Phe Thr 135 Glu Arg Ala Thr Glu Glu Gly Ser Gly Ser Phe Thr Gln Ser Arg Ser 150 155 Ser His Arg Val Ala Arg Val Ala Arg Val Cys Lys Gly Asp Leu Gly 170 165 Gly Lys Lys Ile Leu Gln Lys Lys Trp Thr Ser Phe Leu Lys Ala Arg 180 185 Leu Ile Cys His Ile Pro Leu Tyr Glu Thr Leu Arg Gly Val Cys Ser 200 205 Leu Asp Ala Glu Thr Ser Ser Arg Thr His Phe Tyr Ala Ala Phe Thr 215 Leu Ser Thr Gln Trp Lys Thr Leu Glu Ala Ser Ala Ile Cys Arg Tyr 235 230 Asp Leu Ala Glu Ile Gln Ala Val Phe Ala Gly Pro Tyr Met Glu Tyr 250 245 Gln Asp Gly Ser Arg Arg Trp Gly Arg Tyr Glu Gly Gly Val Pro Glu 265 Pro Arg Pro Gly Ser Cys Ile Thr Asp Ser Leu Arg Ser Gln Gly Tyr 280 Asn Ser Ser Gln Asp Leu Pro Ser Leu Val Leu Asp Phe Val Lys Leu 295 300 His Pro Leu Met Ala Arg Pro Val Val Pro Thr Arg Gly Arg Pro Leu 315 310 Leu Leu Lys Arg Asn Ile Arg Tyr Thr His Leu Thr Gly Thr Pro Val 330 325 Thr Thr Pro Ala Gly Pro Thr Tyr Asp Leu Leu Phe Leu Gly Thr Ala 345 Asp Gly Trp Ile His Lys Ala Val Val Leu Gly Ser Gly Met His Ile 360 Ile Glu Glu Thr Gln Val Phe Arg Glu Ser Gln Ser Val Glu Asn Leu 375 380 Val Ile Ser Leu Leu Gln His Ser Leu Tyr Val Gly Ala Pro Ser Gly 395 390 Val Ile Gln Leu Pro Leu Ser Ser Cys Ser Arg Tyr Arg Ser Cys Tyr 405 410 Asp Cys Ile Leu Ala Arg Asp Pro Tyr Cys Gly Trp Asp Pro Gly Thr 425 430 420 His Ala Cys Ala Ala Ala Thr Thr Ile Ala Asn Arg Thr Ala Leu Ile 440 Gln Asp Ile Glu Arg Gly Asn Arg Gly Cys Glu Ser Ser Arg Asp Thr 455 460 Gly Pro Pro Pro Pro Leu Lys Thr Arg Ser Val Leu Arg Gly Asp Asp 470 475

```
Val Leu Leu Pro Cys Asp Gln Pro Ser Asn Leu Ala Arg Ala Leu Trp
               485
                                   490
Leu Leu Asn Gly Ser Met Gly Leu Ser Asp Gly Gln Gly Gly Tyr Arg
                               505
           500
Val Gly Val Asp Gly Leu Leu Val Thr Asp Ala Gln Pro Glu His Ser
       515
                          520
                                             525
Gly Asn Tyr Gly Cys Tyr Ala Glu Glu Asn Gly Leu Arg Thr Leu Leu
                       535
                                          540
Ala Ser Tyr Ser Leu Thr Val Arg Pro Ala Thr Pro Ala Pro Ala Pro
                   550
                                       555
545
Lys Ala Pro Ala Thr Pro Gly Ala Gln Leu Ala Pro Asp Val Arg Leu
                                   570
               565
Leu Tyr Val Leu Ala Ile Ala Ala Leu Gly Gly Leu Cys Leu Ile Leu
                               585
           580
Ala Ser Ser Leu Leu Tyr Val Ala Cys Leu Arg Glu Gly Arg Arg Gly
                           600
                                               605
Arg Arg Arg Lys Tyr Ser Leu Gly Arg Ala Ser Arg Ala Gly Gly Ser
                       615
                                           620
Ala Val Gln Leu Gln Thr Val Ser Gly Arg Ala Leu Gln Val His Met
625
                   630
                                       635
Gly Ser Met Ser Pro Pro Ser Ala Trp Pro Cys Val Leu Asp Gly Pro
                                  650 655
               645
Glu Thr Arg Gln Val Leu Cys Gln Pro Pro Lys Pro Cys Val His Ser
                                                  670
                               665
His Ala His Met Glu Glu Cys Leu Ser Ala Gly Leu Gln Cys Pro His
                                               685
        675
                           680
Pro His Leu Leu Val His Ser Cys Phe Ile Pro Ala Ser Gly Leu
                       695
                                           700
Gly Val Pro Ser Gln Leu Pro His Pro Ile Trp Ser Ser Pro Ala
                   710
                                       715
Pro Cys Gly Asp Leu Phe Val Lys Ser Leu Gly Thr Gly Gln Pro Gly
                                   730
               725
Glu Val Arg Leu His His Ser Pro Pro Leu Pro Ser Cys Val Ala Leu
           740
                              745
Val Asn Gln Pro Pro His Ser Pro Trp Ser Phe Ser Arg Val
                           760
        755
```

<211> 2951

<212> DNA

<213> homo sapiens

<400> 50

```
60
ccgcggagcc gcgcgtcgct gtctctatgg ccccggatcc gagcgcaaag aaaacggacc
                                                                       120
tcagaaaacc aggactagct ctactgtcgg gggcagggtg accccatcag taacctacaa
                                                                       180
cccctctaga acttcacaac tccctctcac catggagttt gcatttgatg cagaaaggca
tgtgatccct ccctccttct gacctcttag ctggggattc catggccaca caaccctgtg
                                                                       240
                                                                       300
actocatgto coccegatto caggaccoco catggcocca tgattocttg actoctatga
                                                                       360
cettatgace cetgacette caagtgactt cettggactt tgacecetgt gactgtgett
                                                                       420
cccattcccc gccccacaa cctgtgactc tggctccctt tgggggtctt gttagtctgg
                                                                       480
gcctccccag gaagatgtgg gggaggctct ggccctcct cctcagcatc ctcacagcaa
                                                                       540
ctgcagtccc aggaccctca ctgcggagac cgtctagaga actagatgcc acccctcgga
                                                                       600
tgaccatacc ctatgaagag ctctctggga cccggcactt caagggccaa gcccagaact
actcaacact gctgctggag gaggcctcag caaggctgct ggtgggagcc cgaggtgccc
                                                                       660
                                                                       720
tgttctctct cagtgccaac gacataggag atggggctca caaagagatc cactgggaag
cctccccaga gatgcaaagc aaatgtcatc aaaaagggaa aaacaaccag acggagtgct
                                                                       780
```

ttaaccatqt	gcggttcctg	cagoggotca	attctaccca	cctctatgca	tgtgggactc	840
	gccctctgt					900
	gaaggagaag					960
	cctctacaca					1020
	cccacactcc					1080
	gttctccgtc					1140
	ctacttcttc					1200
	cagtcaccgt					1260
	cctgcagaag					1320
	tgagacactg					1380
	tgcagccttc					1440
	tgacctggca					1500
	ccggcgctgg					1560
	agattcattg					1620
	ctttgtaaag					1680
	gctgctcaag					1740
	tggacctacc					1800
	agtcctgggc					1860
	tgtggaaaat					1920
	agtcatccag					1980
	ggcccgagac					2040
	catagccaac					2100
	cagcagggat					2160
gagatgatga	tgtcctcctg	ccctgtgacc	agccatccaa	cctggcccgg	gccttgtggc	2220
	gagcatgggc					2280
	tacagatgca					2340
	ccgcaccctg					2400
	aaaagcccct					2460
	agccattgcc					2520
	ctgtctgcgg					2580
	ggcaggagga					2640
	tgagggtgat					2700
	ggagggagcc					2760
	tggcttggtg					2820
	gaggcagagc					2880
	catcctggaa					2940
gctctgtctg						2951